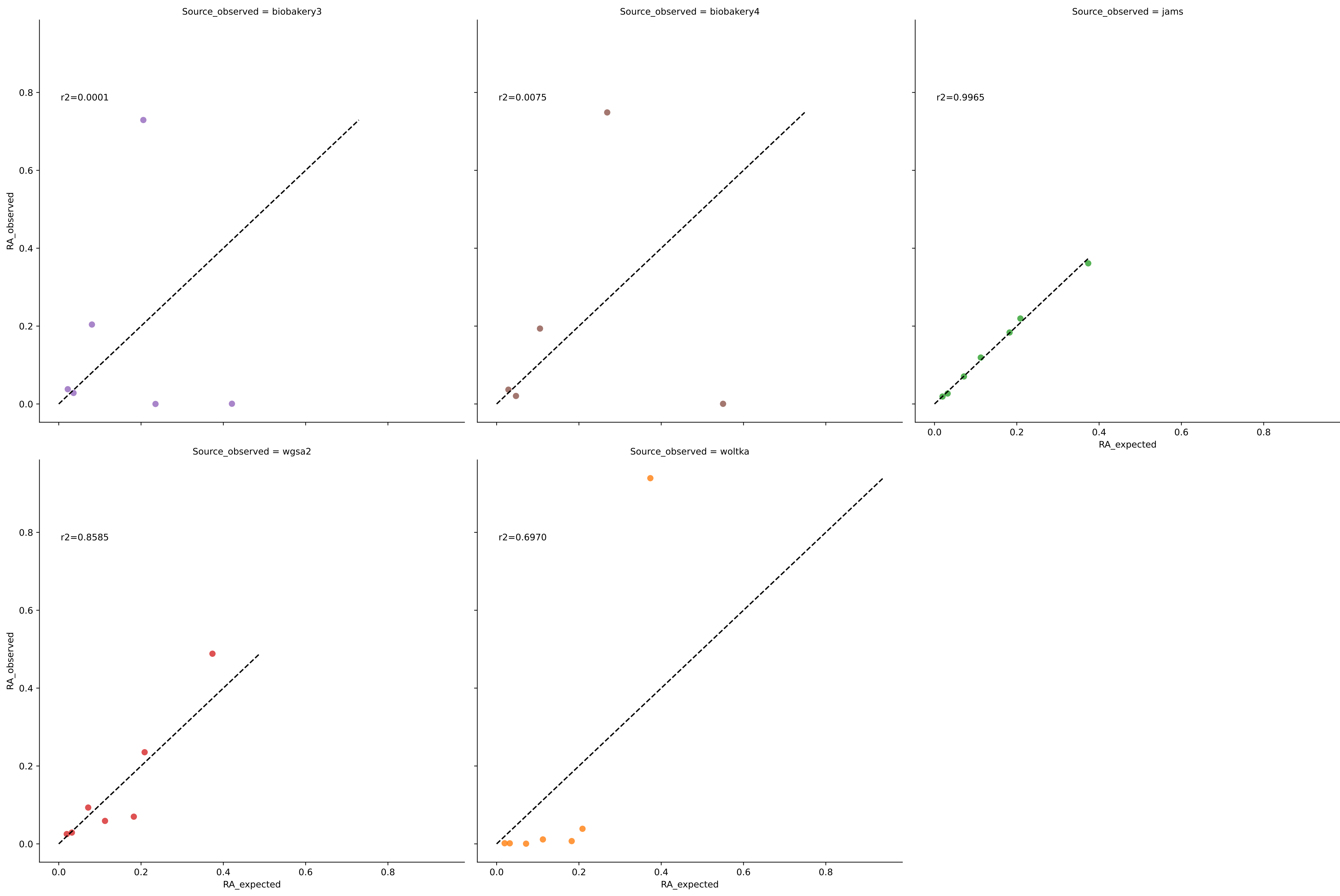
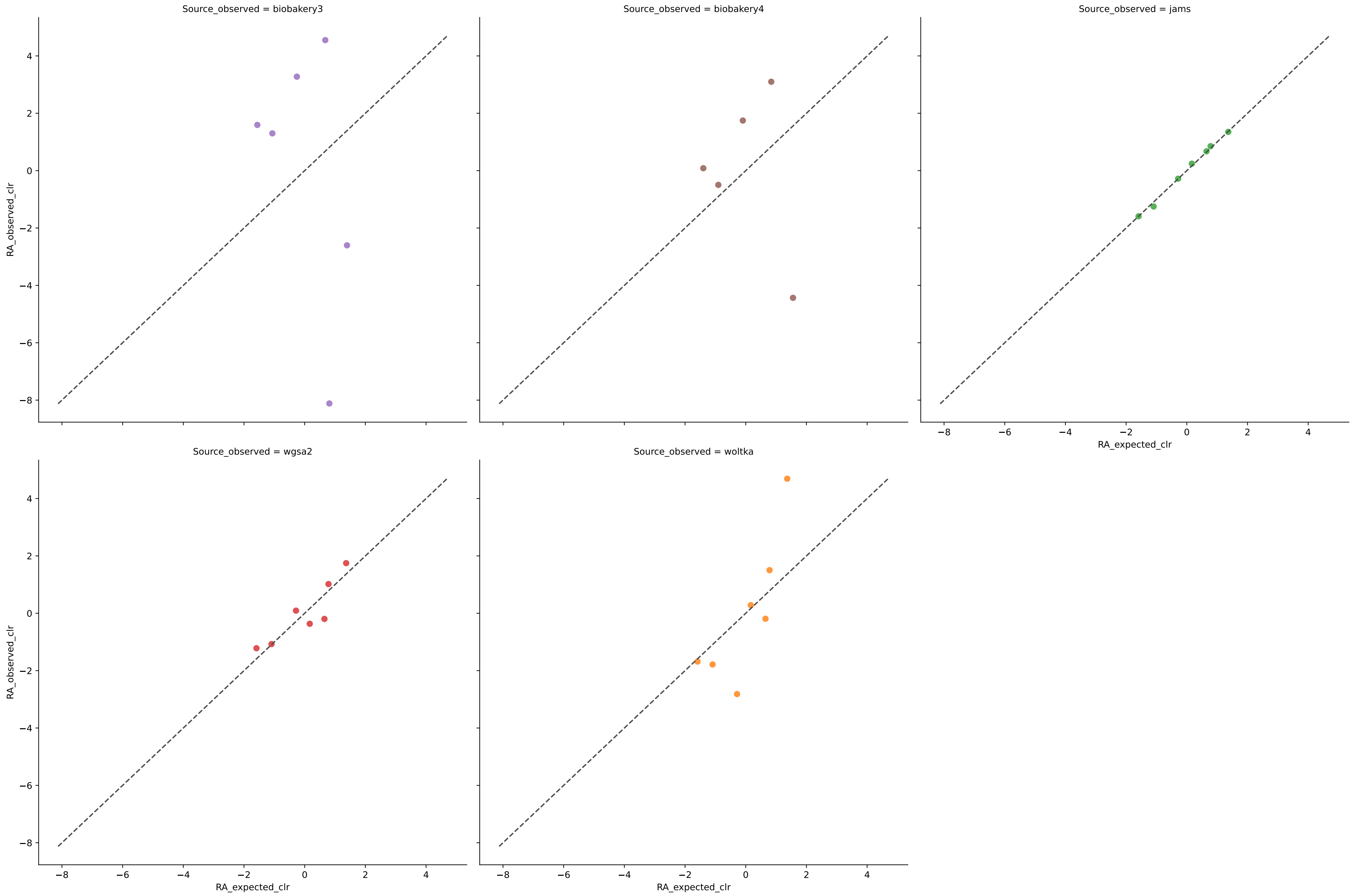


Bivariate Linear Regression for Sample S1 in Experiment bmock12 (Genus at filter threshold 0)

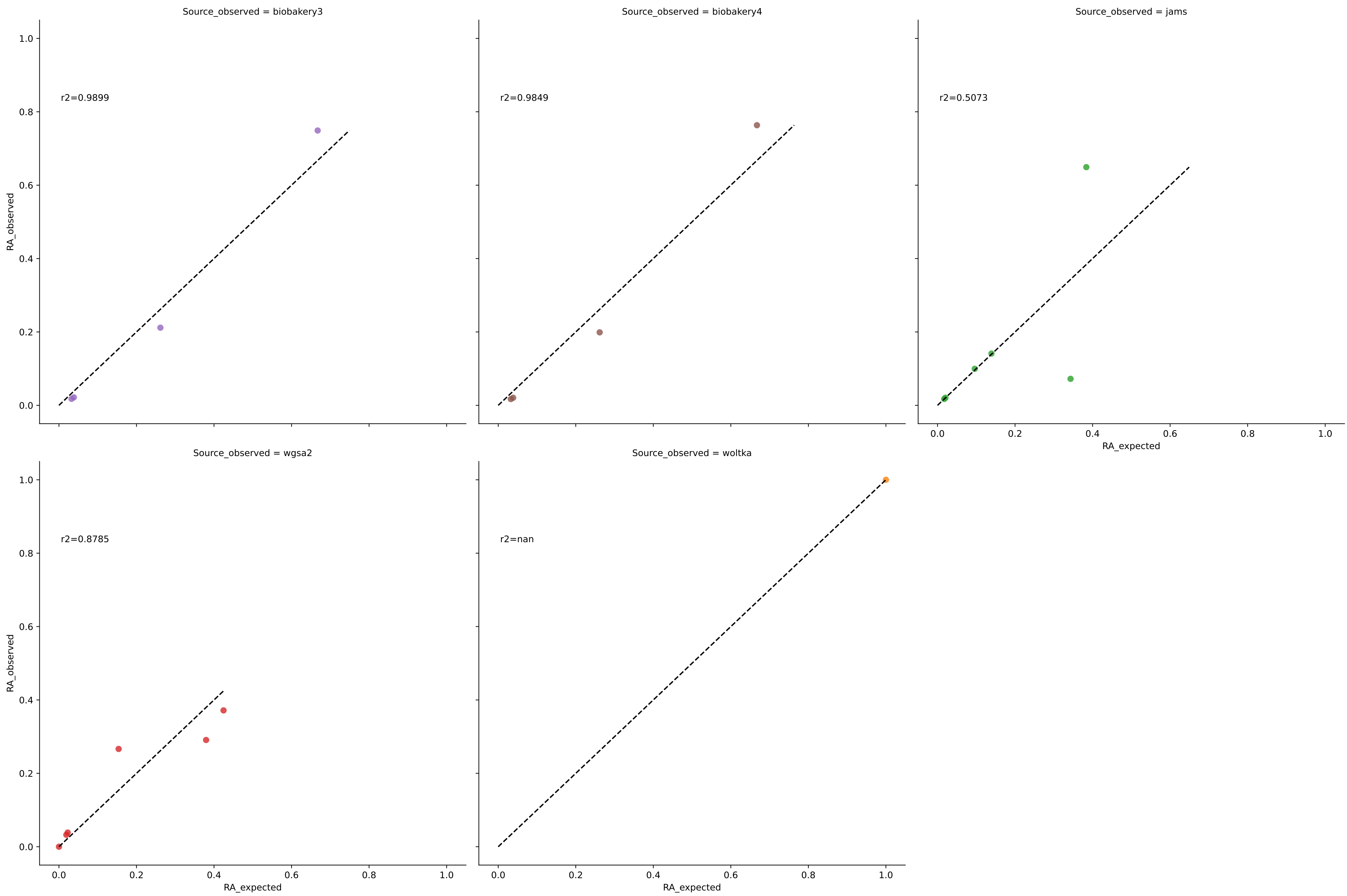


Bivariate Linear Regression for Sample S1 in Experiment bmock12 (Genus at filter threshold 0)

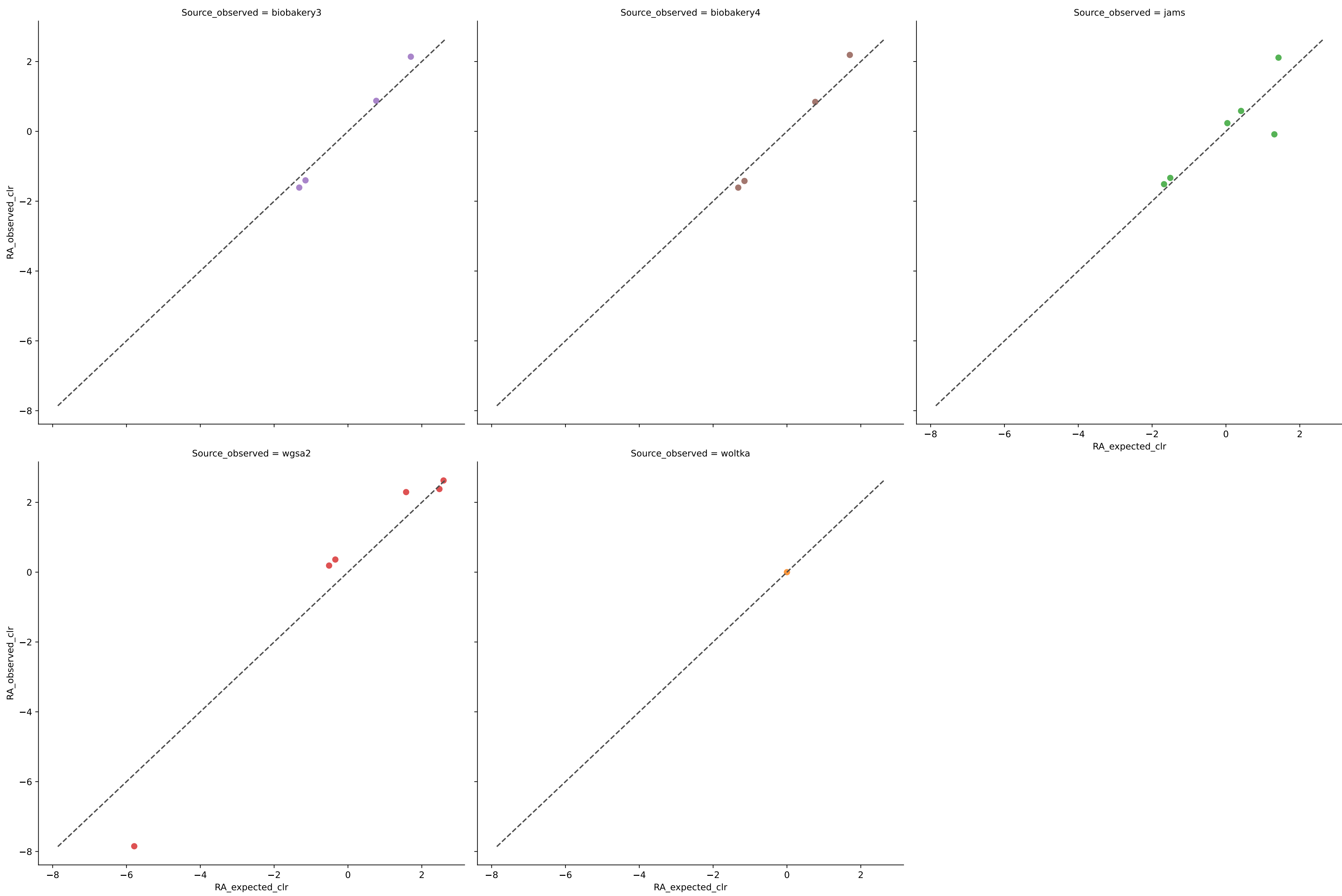


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
biobakery3	6	0.0001	0.2210	11.7793	0.3369	0.2949	100.0000	0.0000
biobakery4	5	0.0075	0.2305	6.8378	0.4237	0.3290	100.0000	0.0000
jams	7	0.9965	0.0053	0.1890	0.9813	0.0070	100.0000	0.0000
wgsa2	7	0.8585	0.0483	1.2171	0.8311	0.0654	100.0000	0.0000
woltka	7	0.6970	0.1616	4.3790	0.4344	0.2379	100.0000	0.0000

Bivariate Linear Regression for Sample S1 in Experiment bmock12 (Species at filter threshold 0)

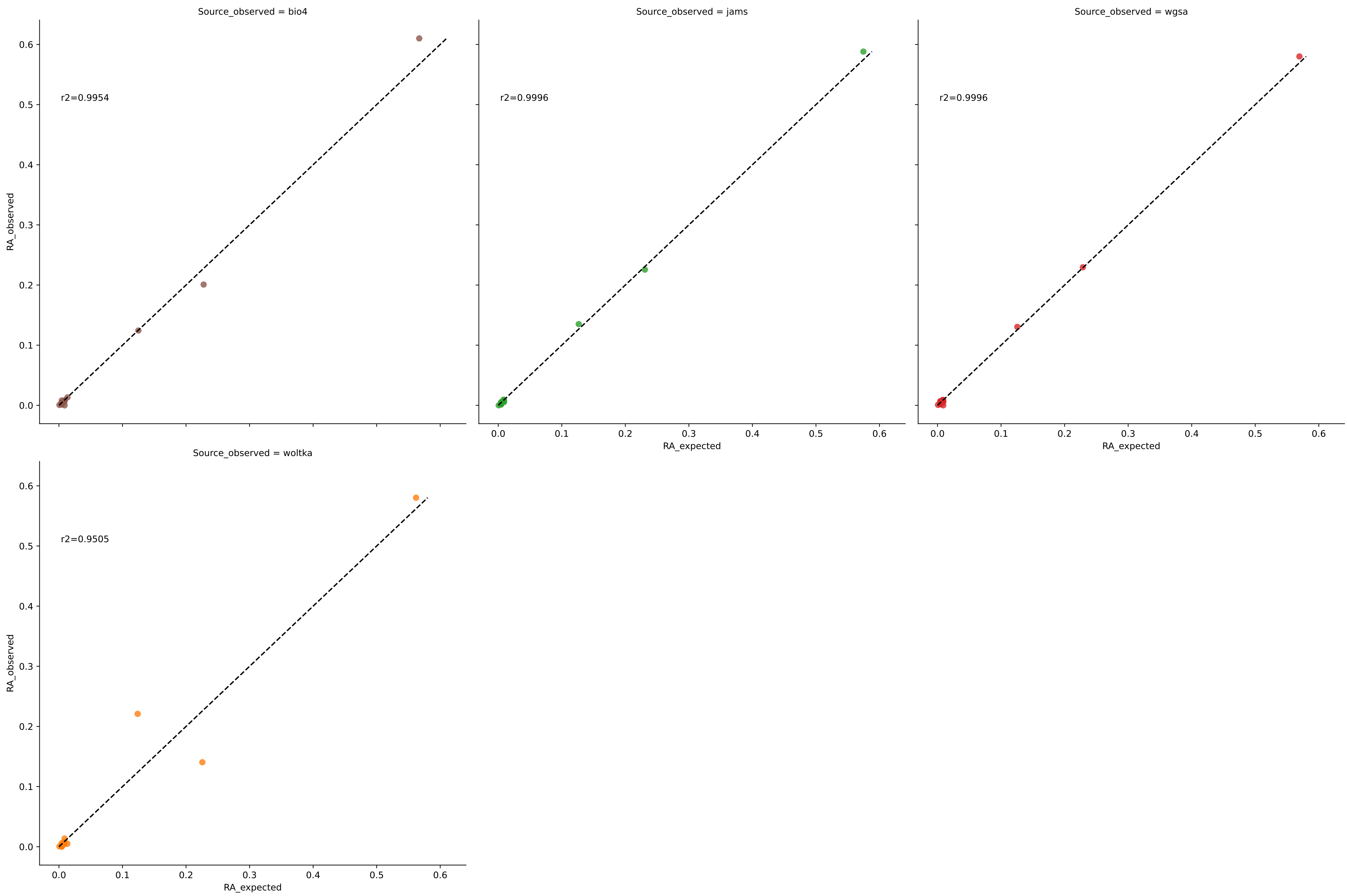


Bivariate Linear Regression for Sample S1 in Experiment bmock12 (Species at filter threshold 0)

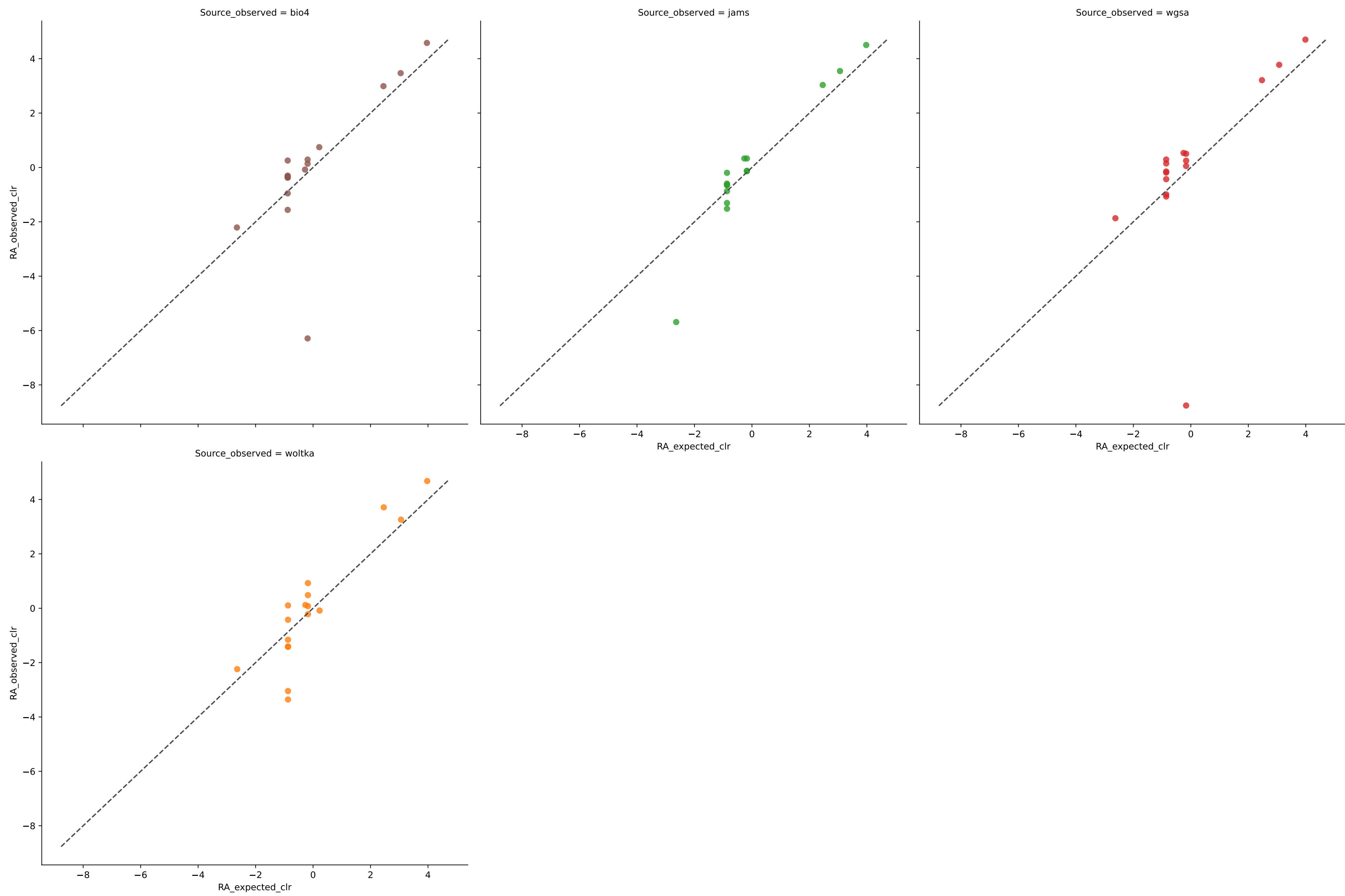


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
biobakery3	4	0.9899	0.0408	0.5907	0.9184	0.0492	100.0000	0.0000
biobakery4	4	0.9849	0.0480	0.6324	0.9040	0.0586	100.0000	0.0000
jams	6	0.5073	0.0903	1.5970	0.7290	0.1548	100.0000	0.0000
wgsa2	6	0.8785	0.0473	2.3991	0.8582	0.0629	100.0000	0.0000
woltka	1	nan	0.0000	0.0000	1.0000	0.0000	100.0000	0.0000

Bivariate Linear Regression for Sample S1 in Experiment camisimGI (Genus at filter threshold 0)

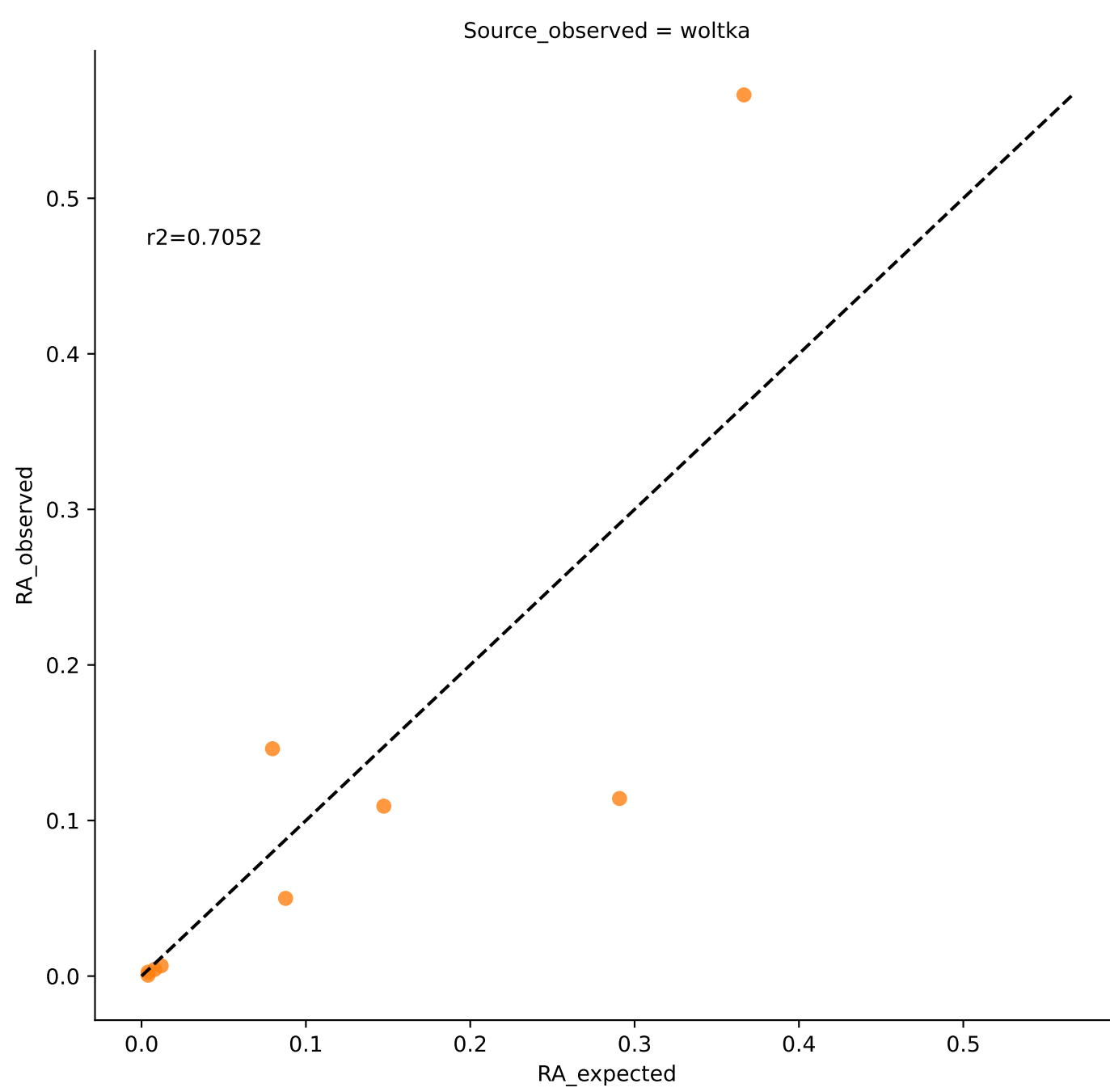
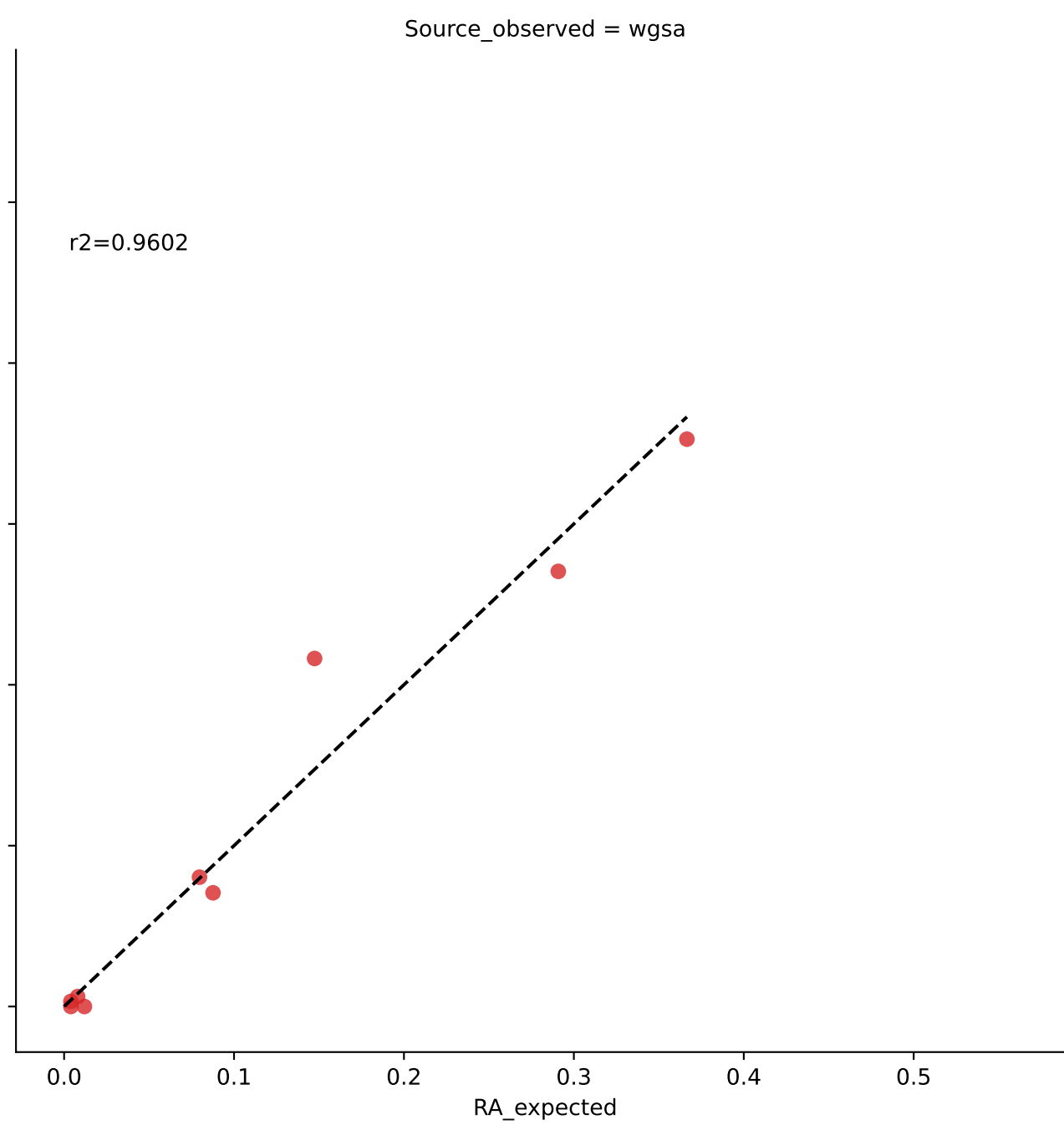
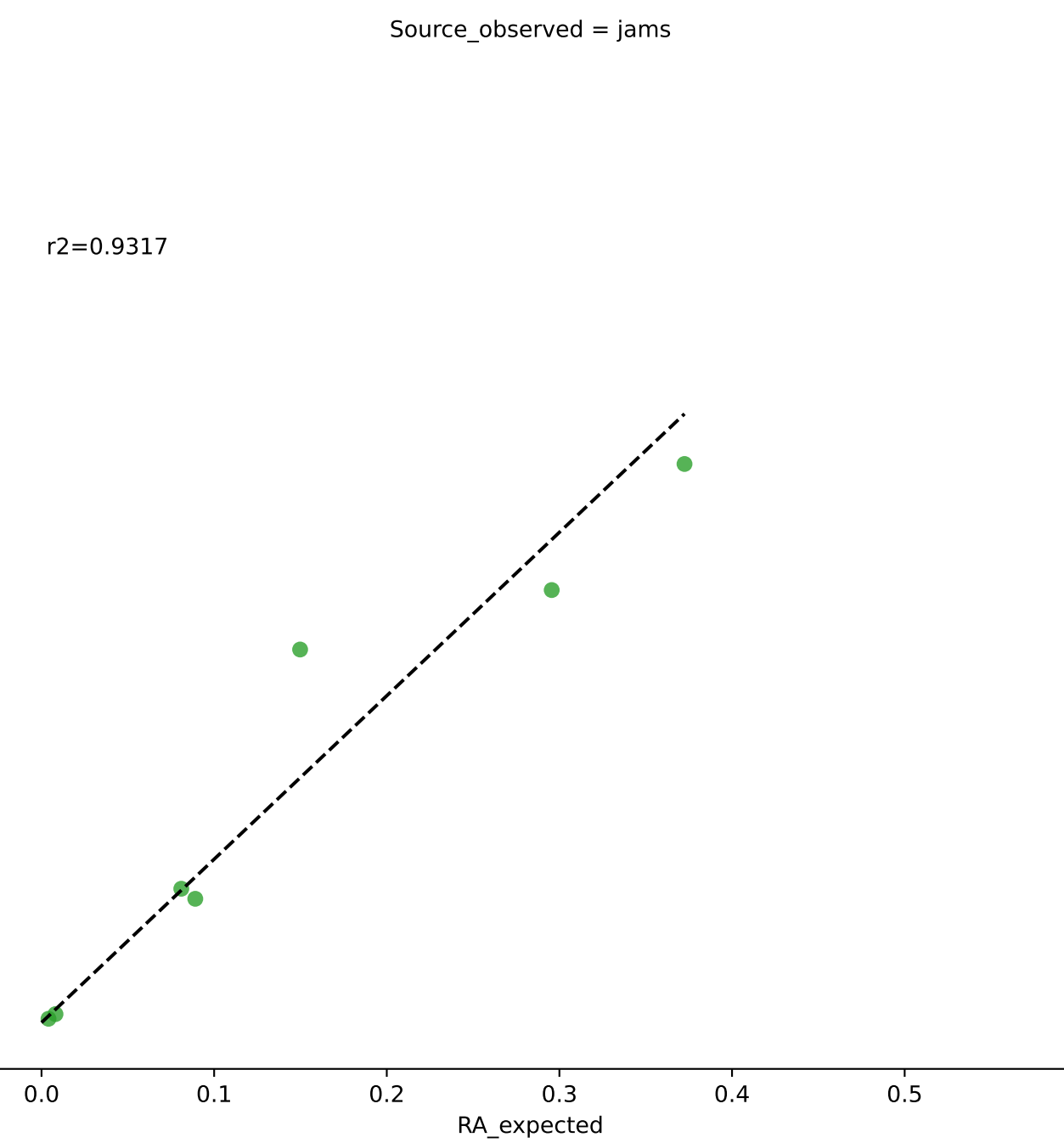
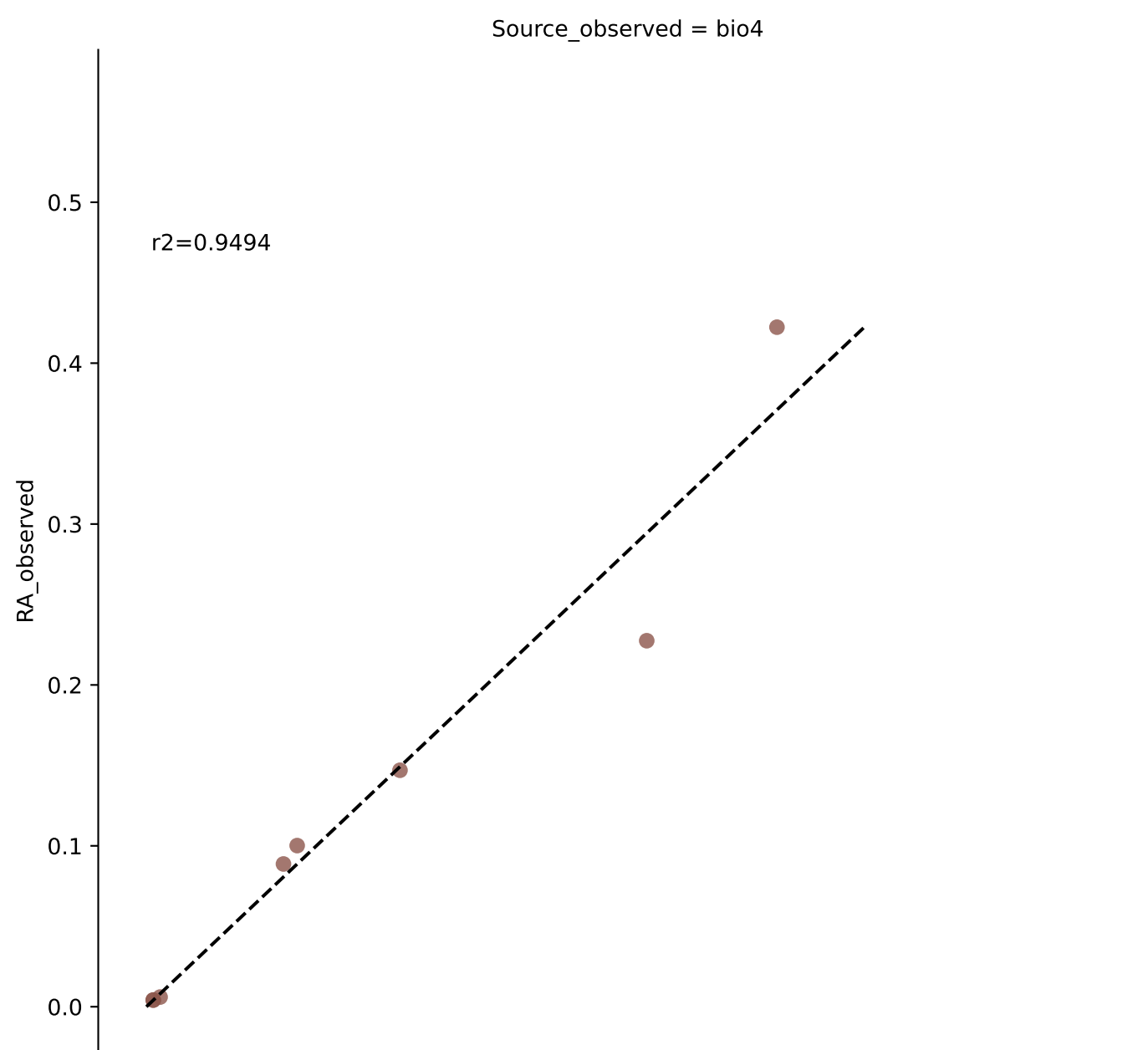


Bivariate Linear Regression for Sample S1 in Experiment camisimGI (Genus at filter threshold 0)

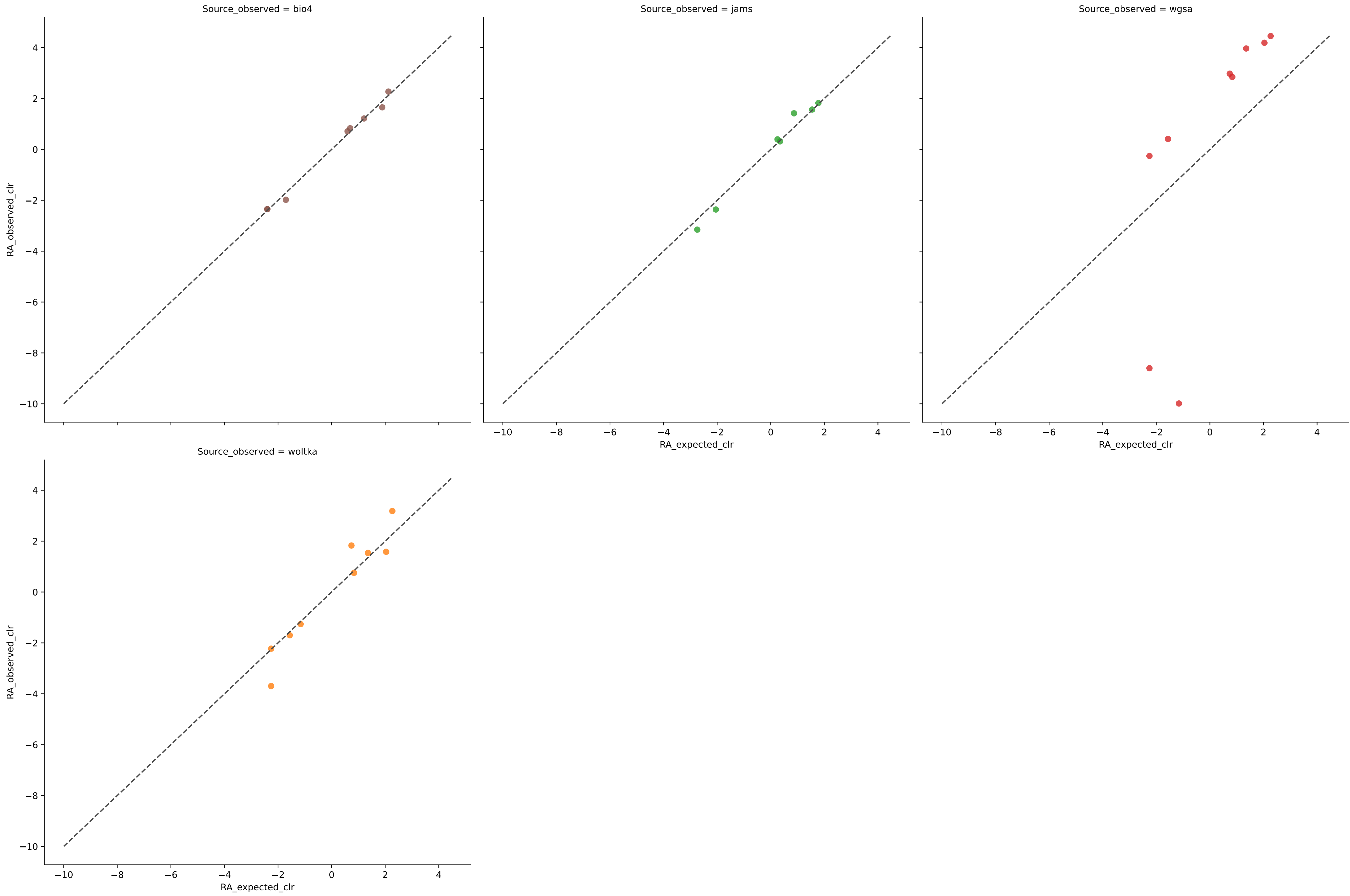


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	16	0.9954	0.0059	6.4683	0.9531	0.0130	100.0000	0.0000
jams	15	0.9996	0.0031	3.4640	0.9766	0.0047	100.0000	0.0000
wgsa	16	0.9996	0.0027	8.9863	0.9787	0.0040	100.0000	0.0000
woltka	17	0.9505	0.0143	4.1172	0.8784	0.0318	100.0000	0.0000

Bivariate Linear Regression for Sample S2 in Experiment camisimGI (Genus at filter threshold 0)

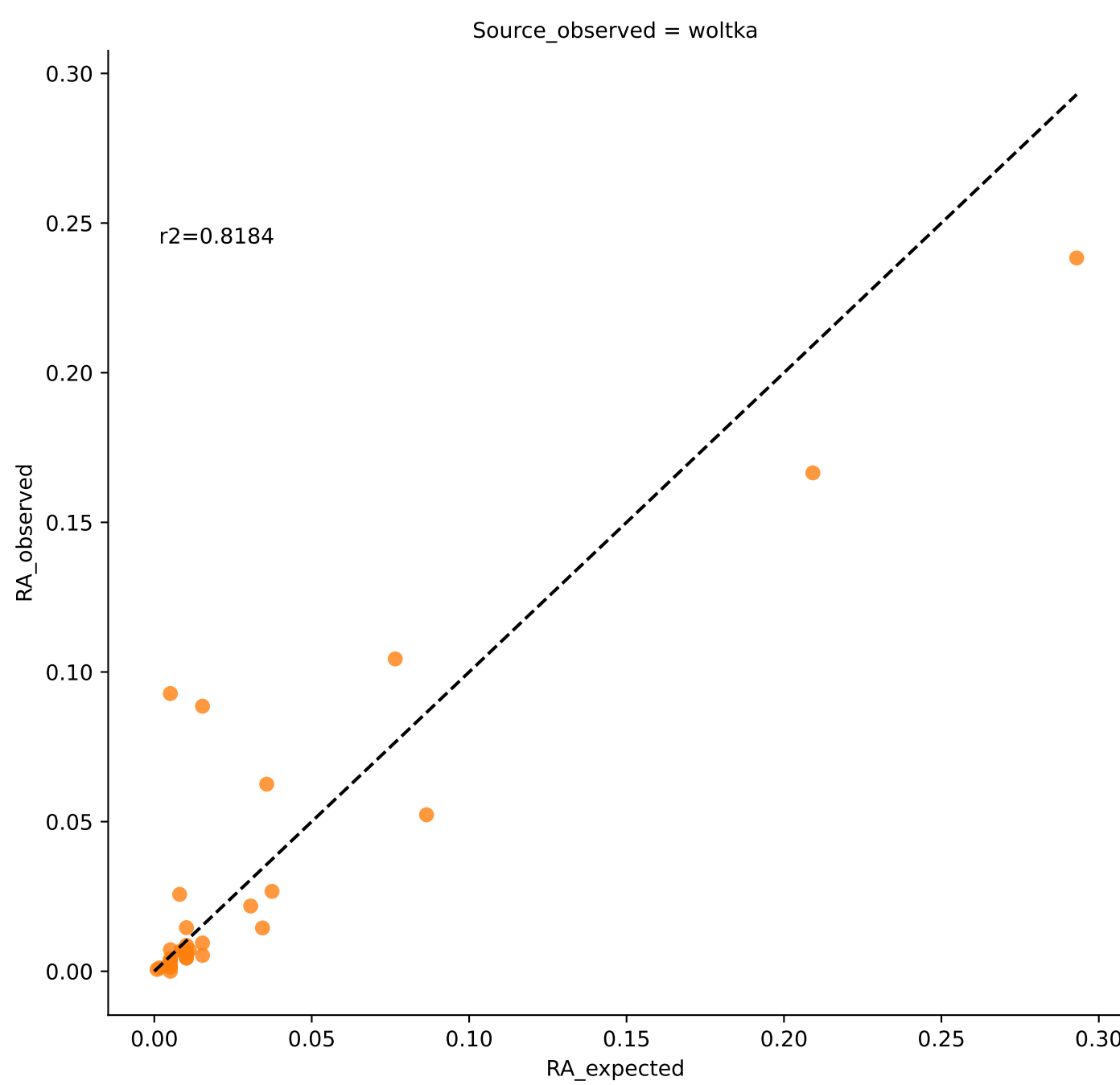
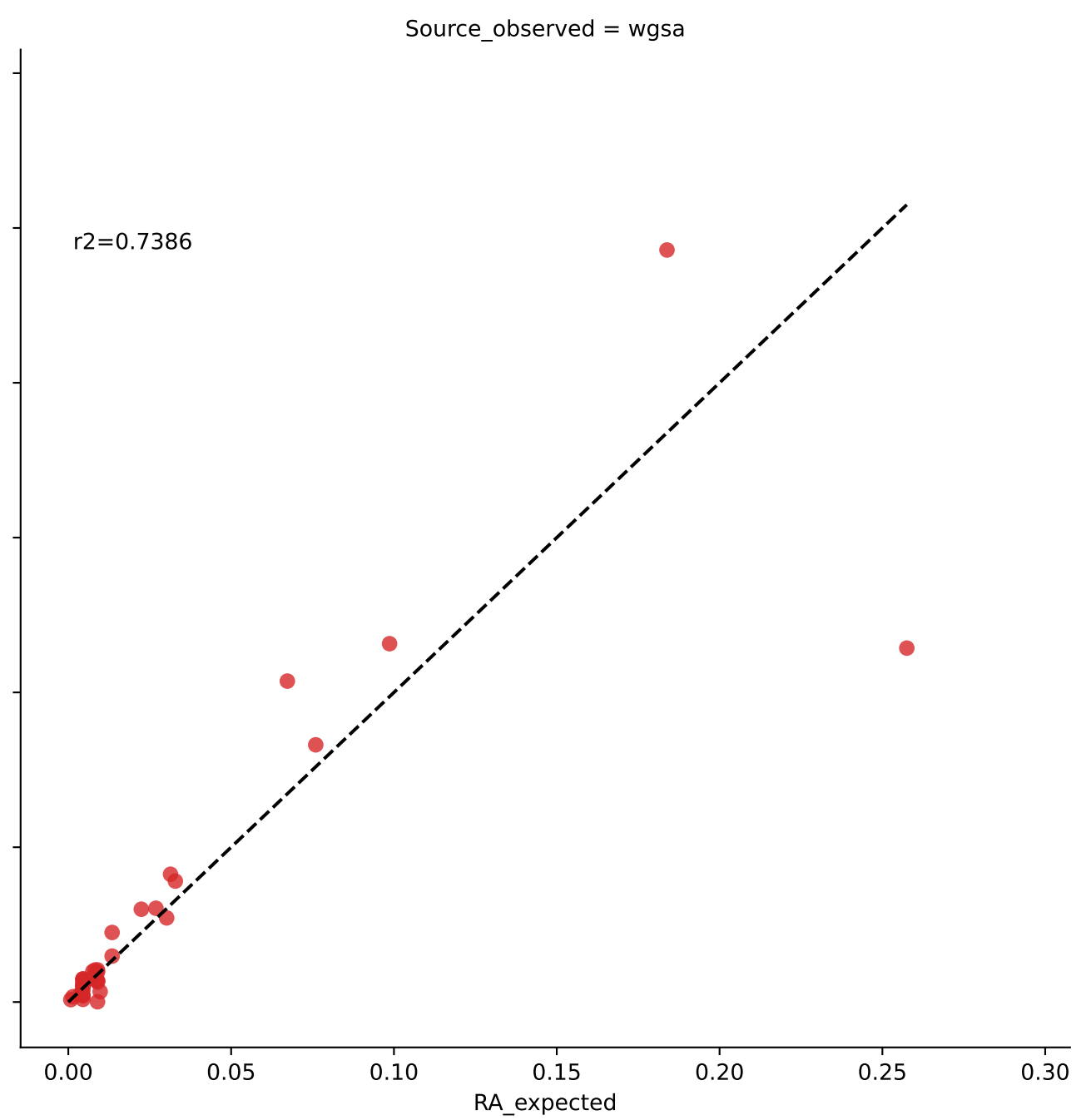
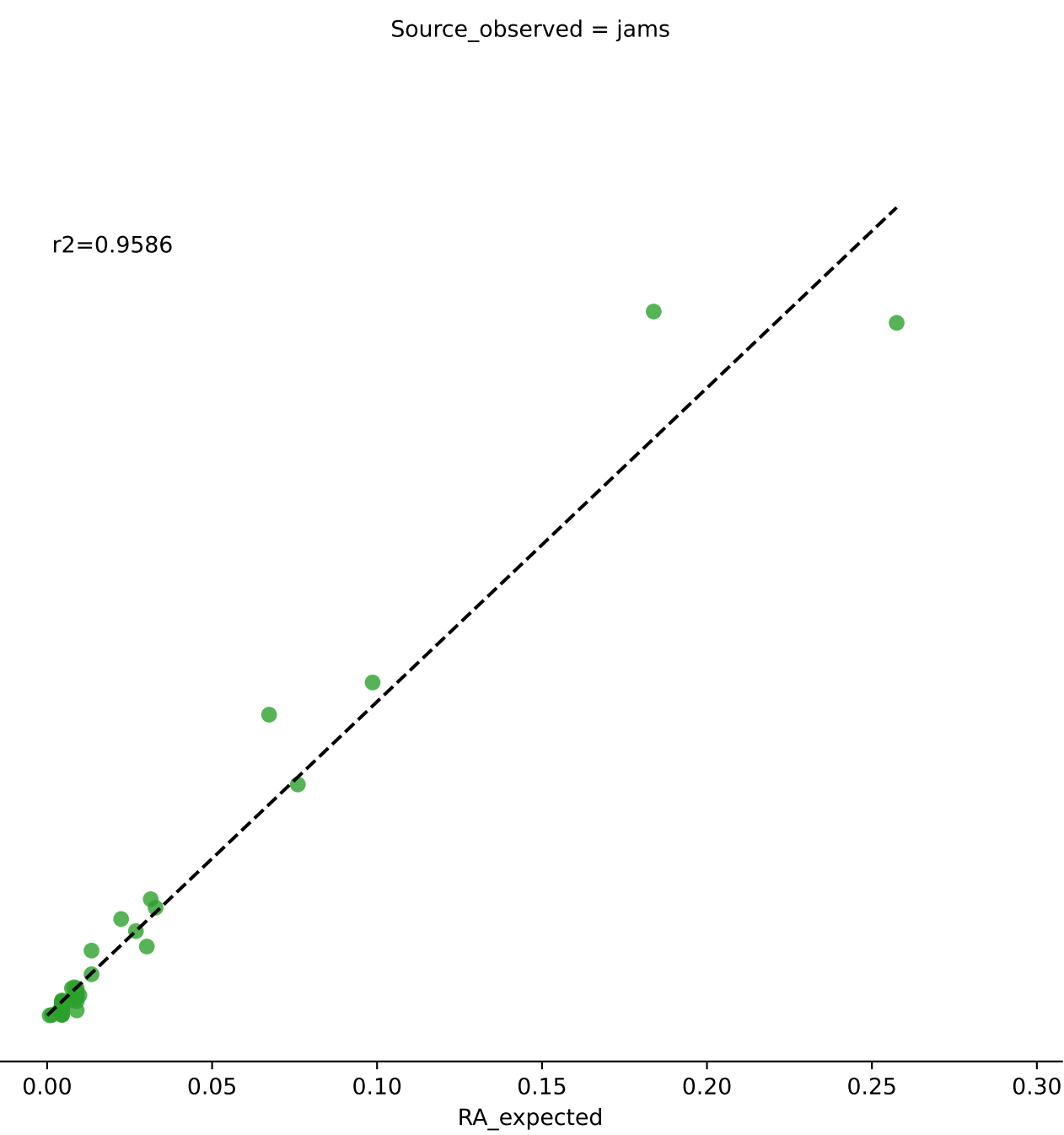
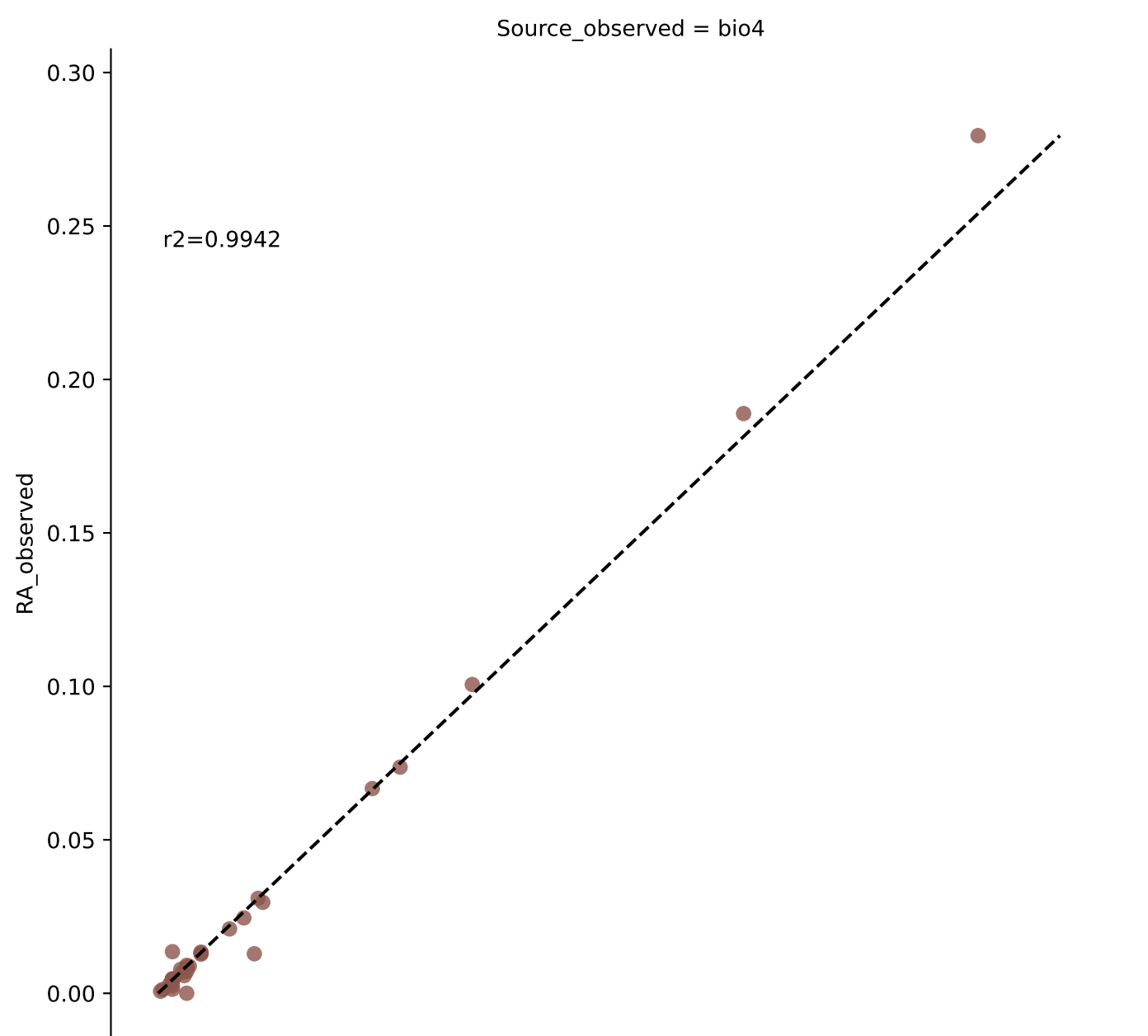


Bivariate Linear Regression for Sample S2 in Experiment camisimGI (Genus at filter threshold 0)

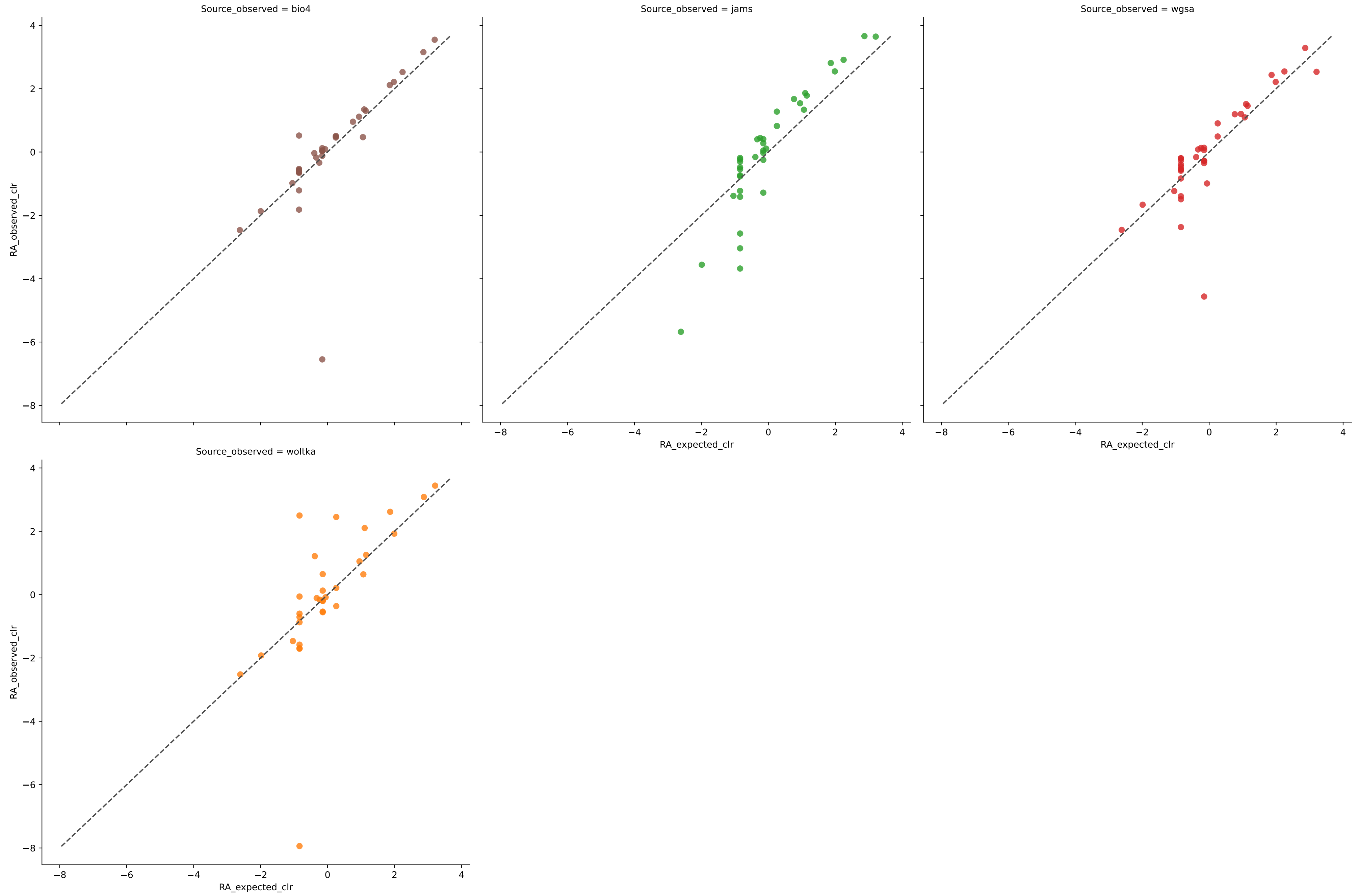


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	8	0.9494	0.0178	0.4385	0.9288	0.0302	100.0000	0.0000
jams	7	0.9317	0.0227	0.7701	0.9206	0.0343	100.0000	0.0000
wgsa	9	0.9602	0.0155	12.3060	0.9303	0.0254	100.0000	0.0000
woltka	9	0.7052	0.0592	2.0925	0.7336	0.0934	100.0000	0.0000

Bivariate Linear Regression for Sample S1 in Experiment camisimGI (Species at filter threshold 0)

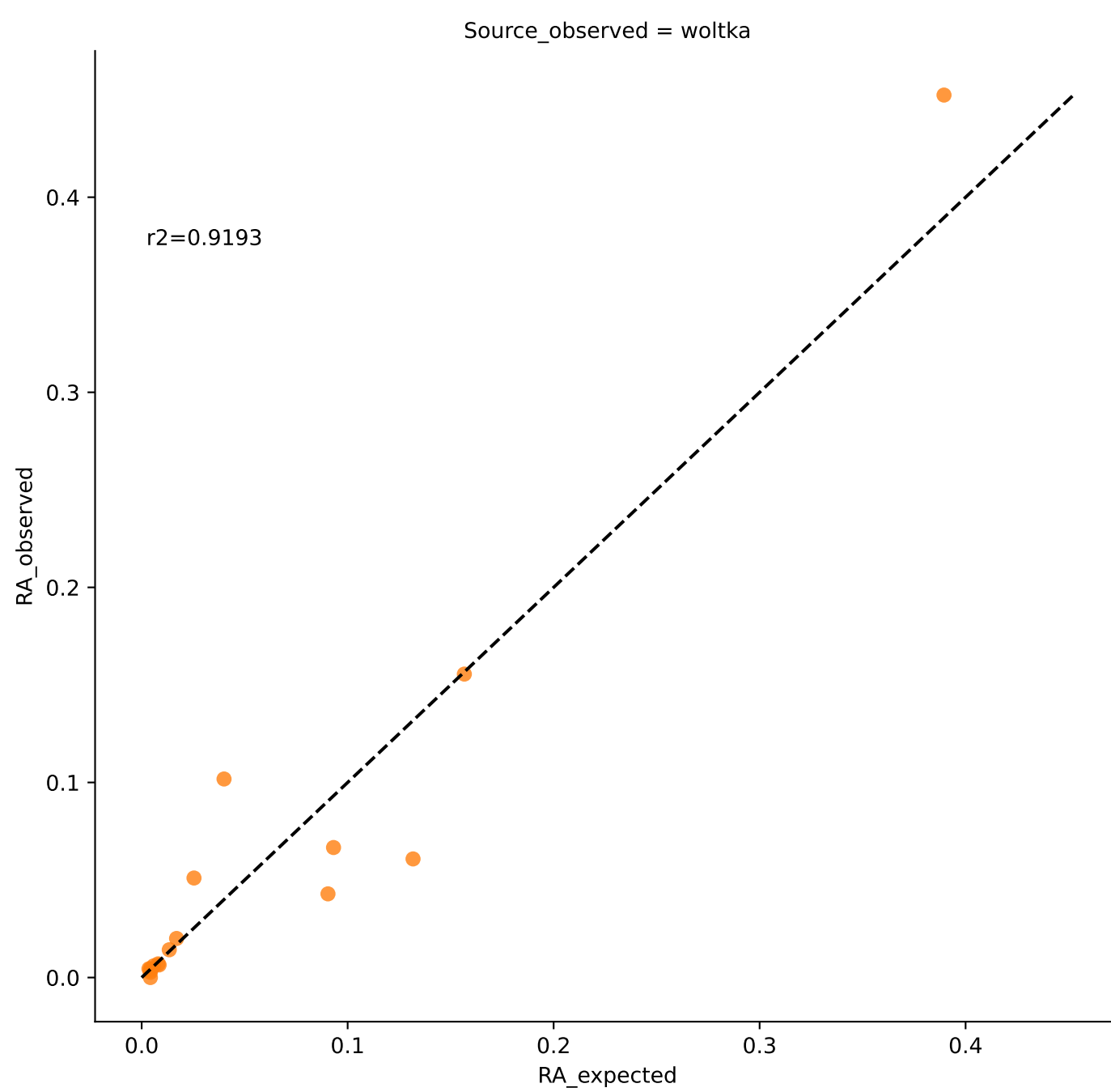
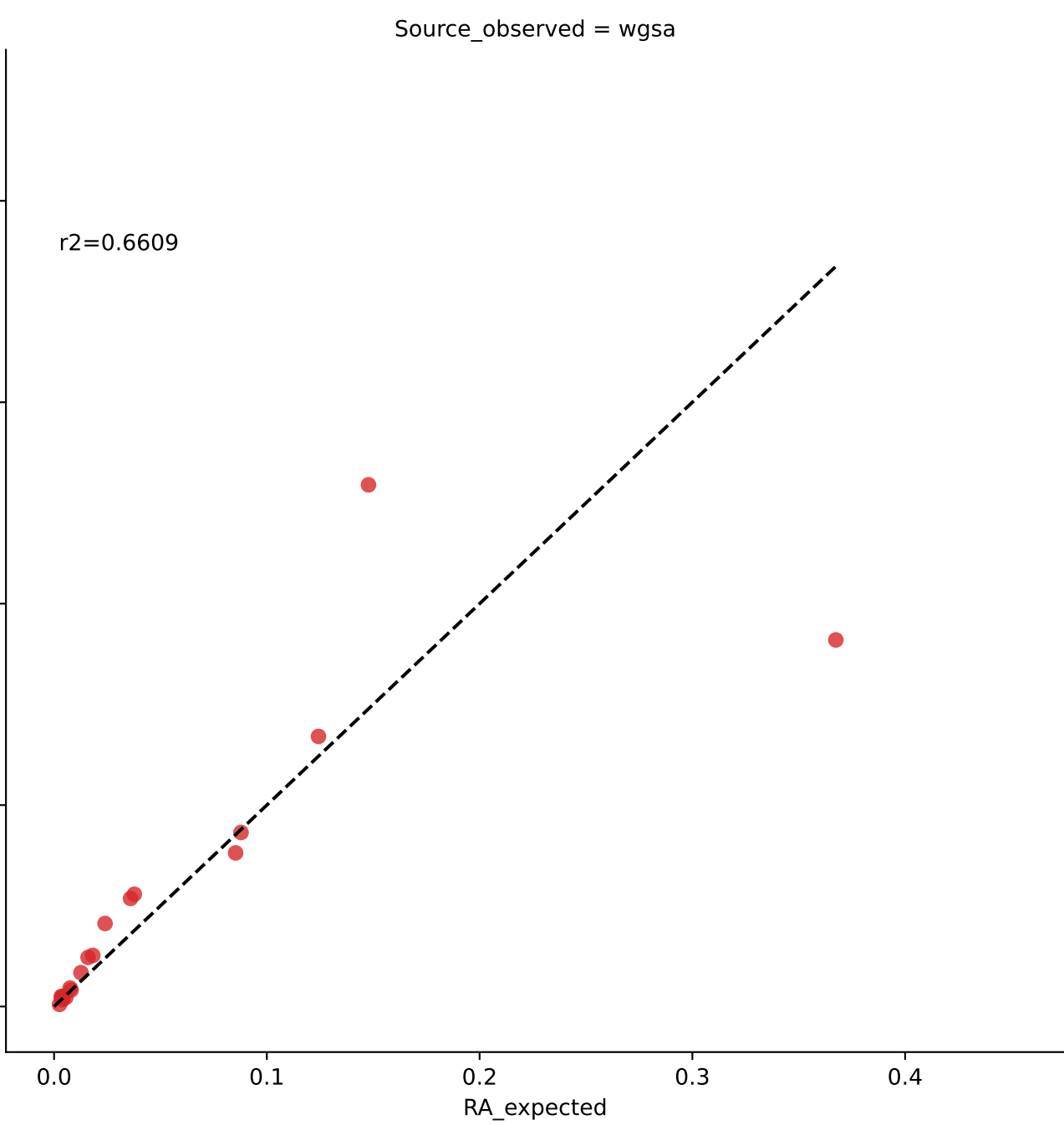
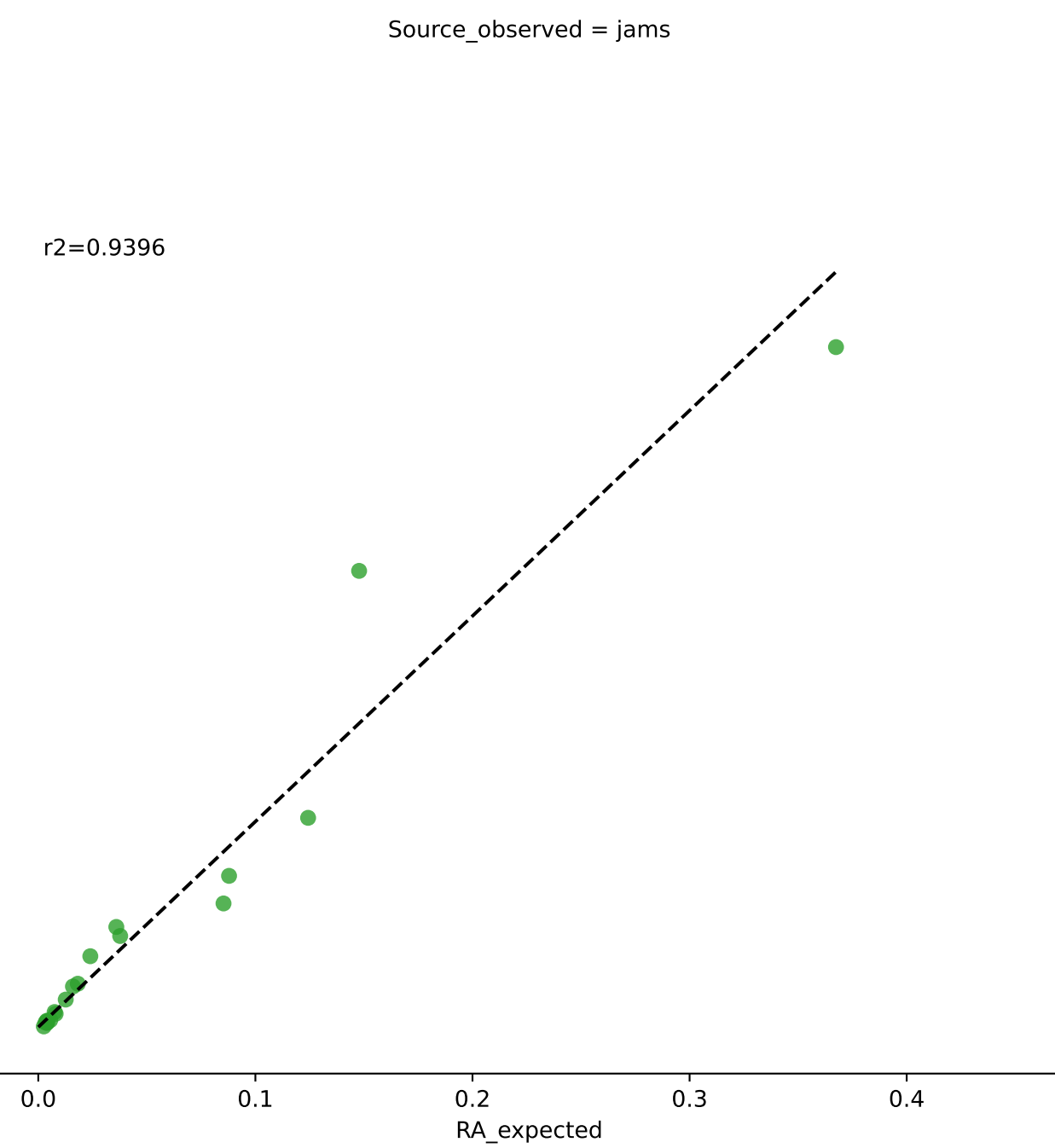
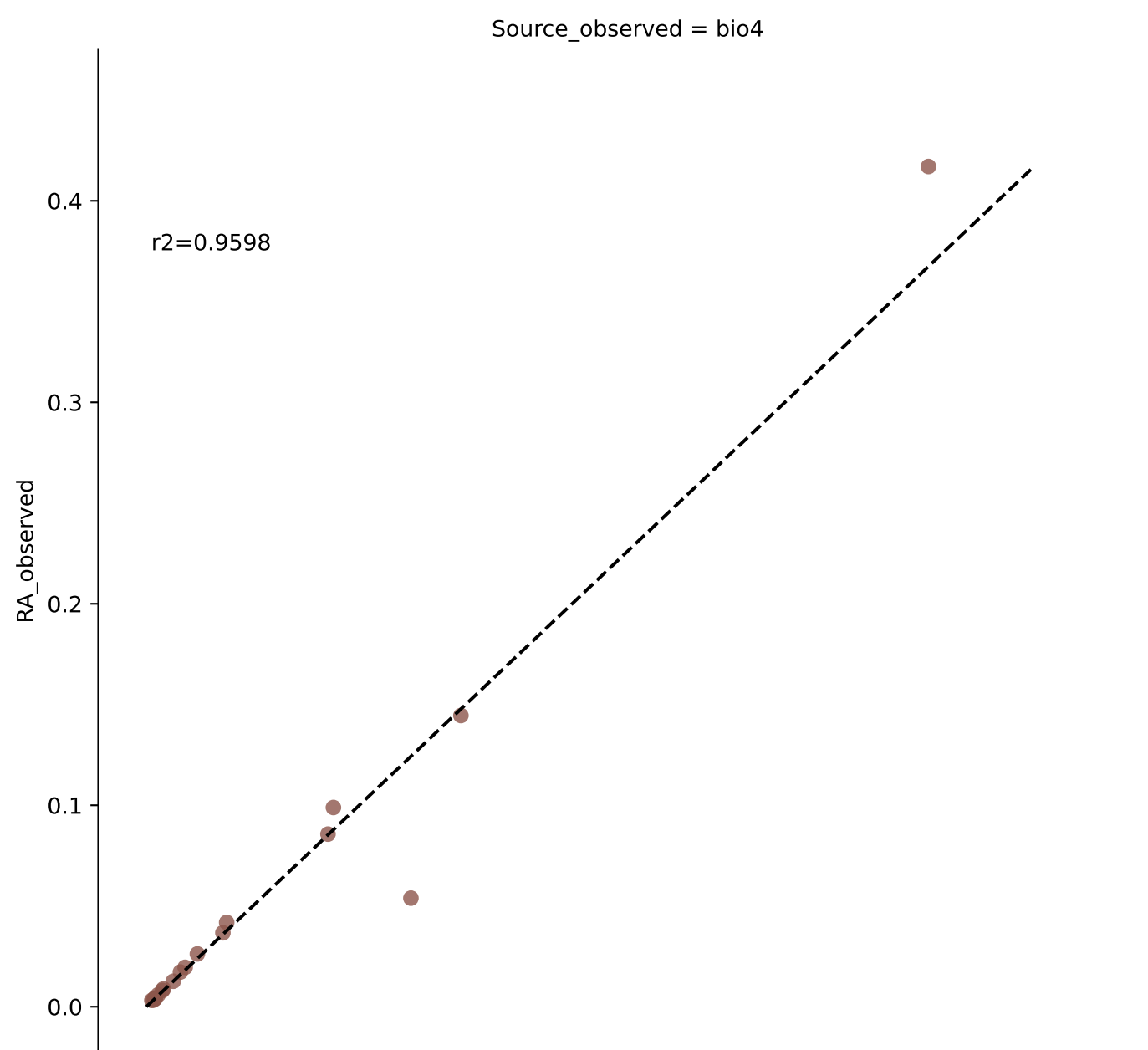


Bivariate Linear Regression for Sample S1 in Experiment camisimGI (Species at filter threshold 0)

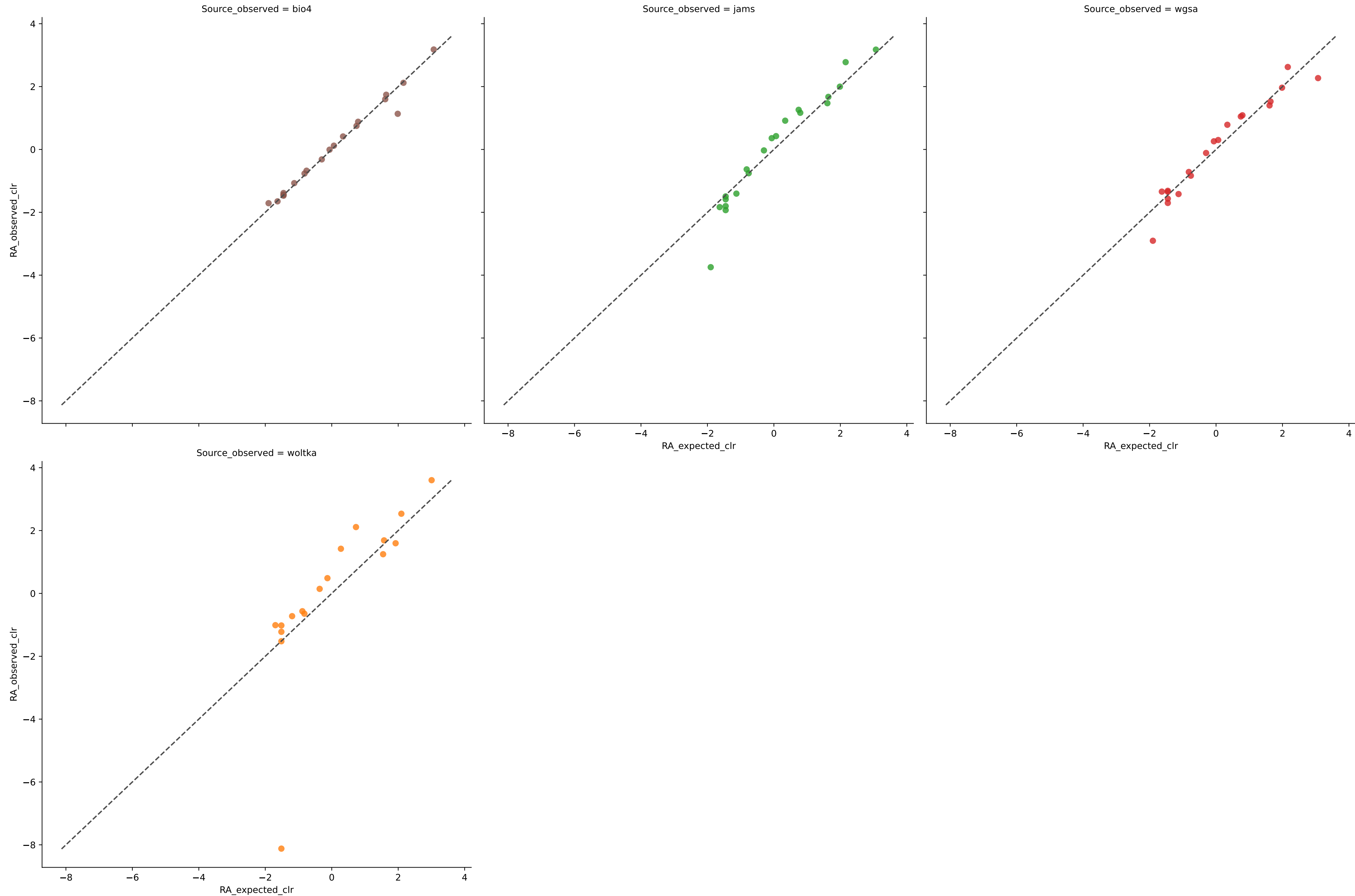


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	38	0.9942	0.0025	6.7698	0.9527	0.0056	100.0000	0.0000
jams	37	0.9586	0.0055	6.2135	0.8985	0.0108	100.0000	0.0000
wgsa	37	0.7386	0.0096	5.2876	0.8219	0.0266	100.0000	0.0000
woltka	33	0.8184	0.0145	8.6684	0.7603	0.0257	100.0000	0.0000

Bivariate Linear Regression for Sample S2 in Experiment camisimGI (Species at filter threshold 0)

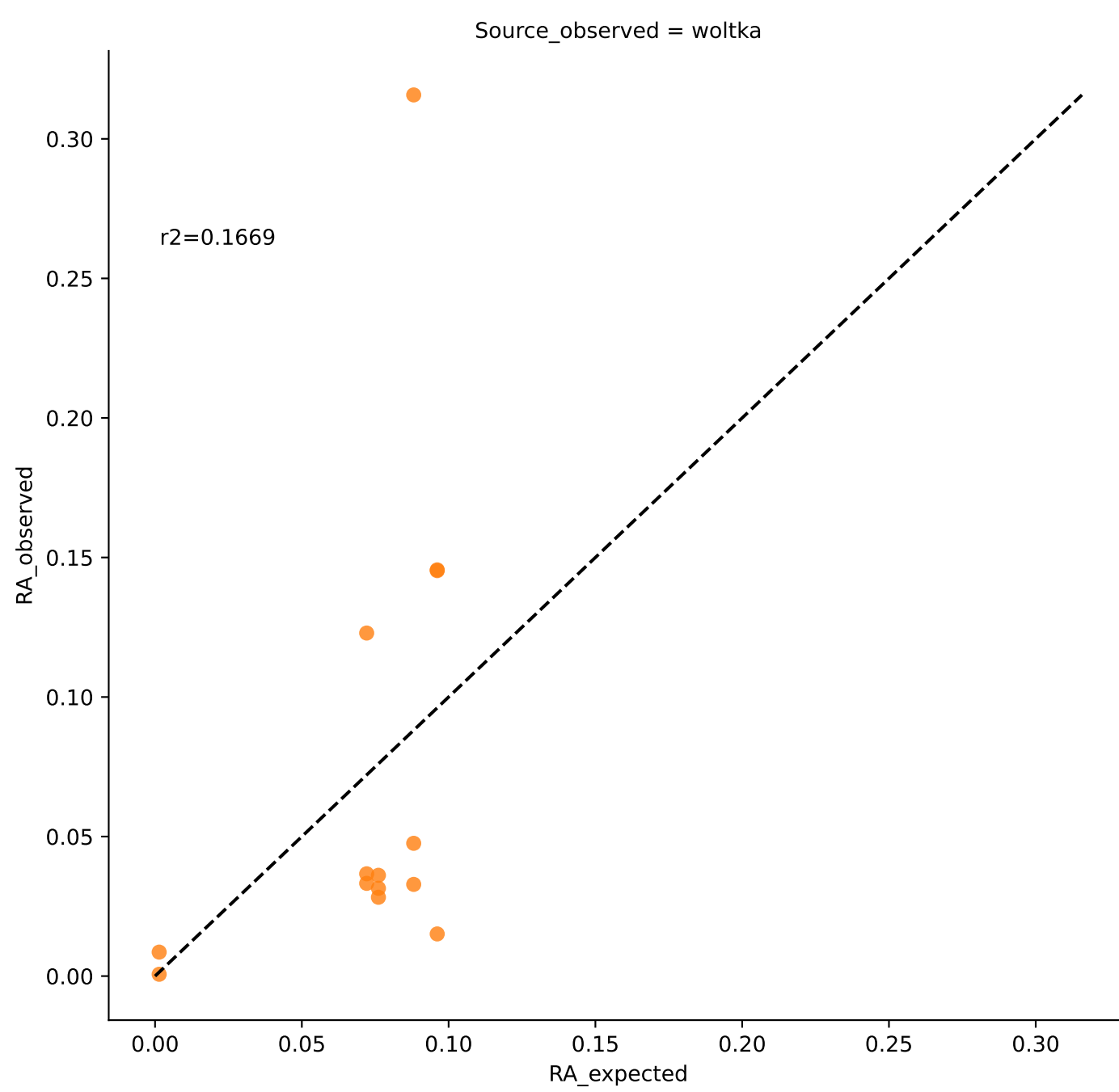
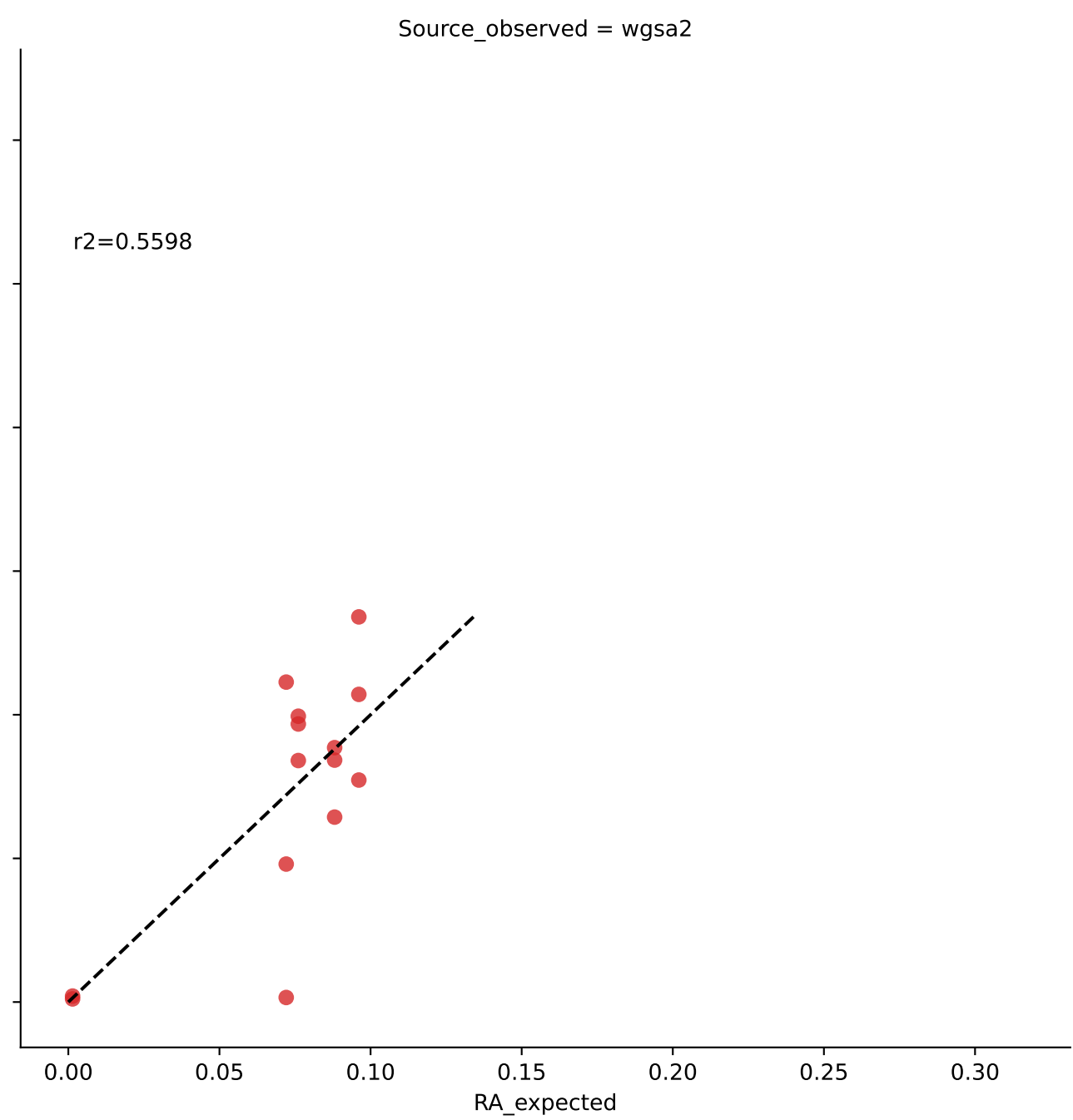
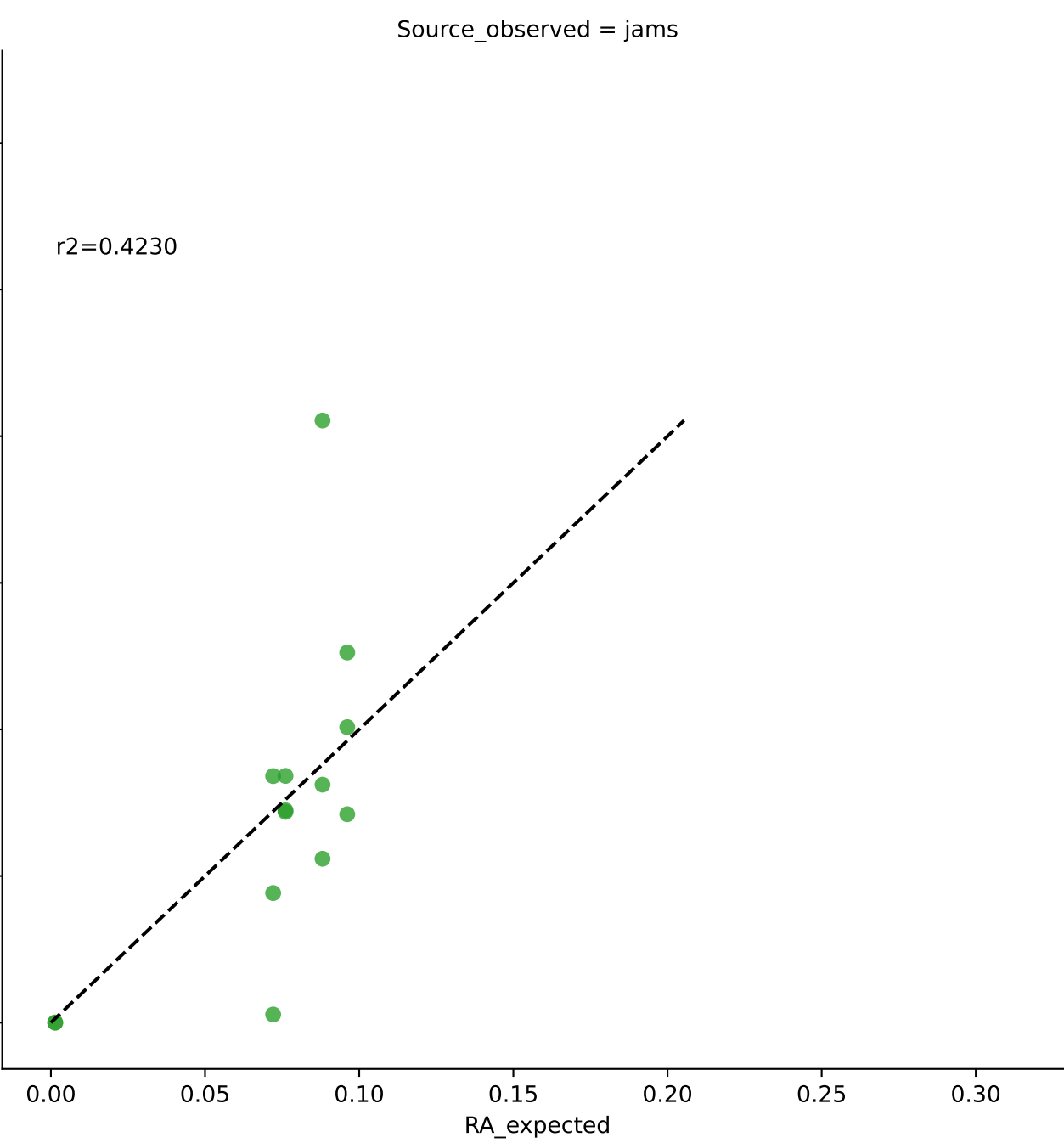
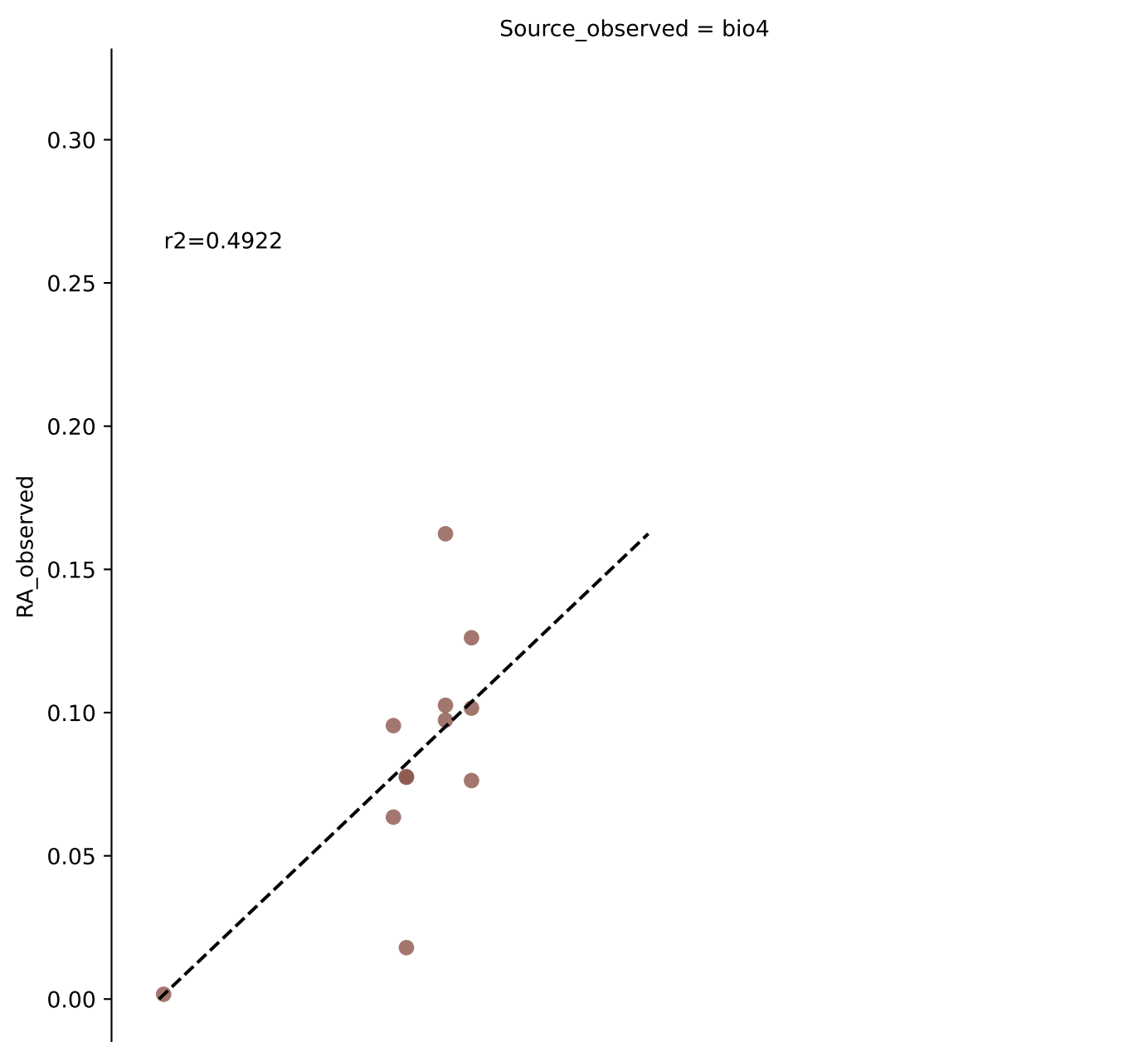


Bivariate Linear Regression for Sample S2 in Experiment camisimGI (Species at filter threshold 0)

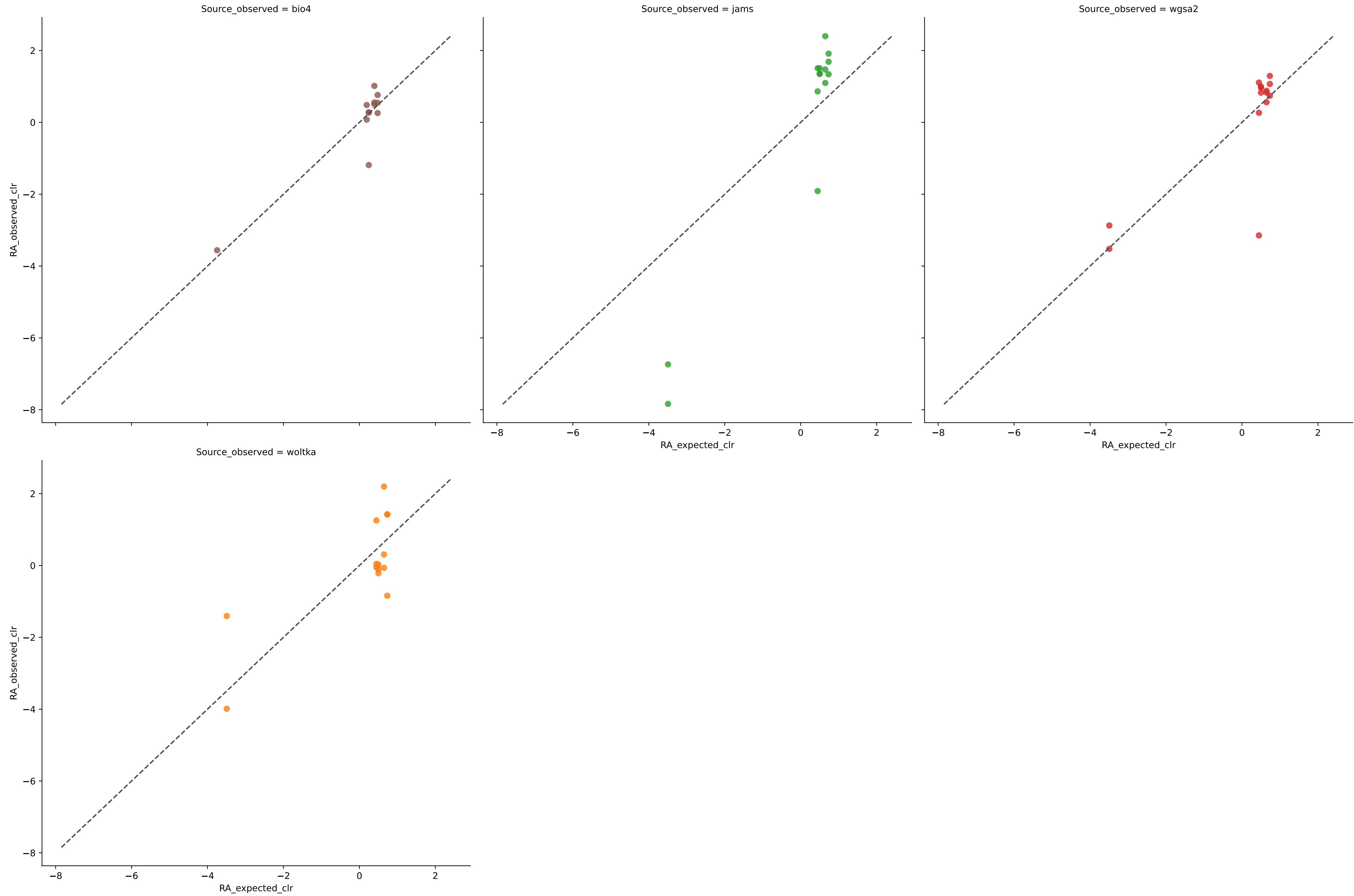


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	20	0.9598	0.0074	0.9077	0.9264	0.0195	100.0000	0.0000
jams	20	0.9396	0.0112	2.3368	0.8885	0.0207	100.0000	0.0000
wgsa	20	0.6609	0.0199	1.6681	0.8009	0.0490	100.0000	0.0000
woltka	17	0.9193	0.0183	7.0278	0.8445	0.0311	100.0000	0.0000

Bivariate Linear Regression for Sample EG in Experiment nist (Genus at filter threshold 0)

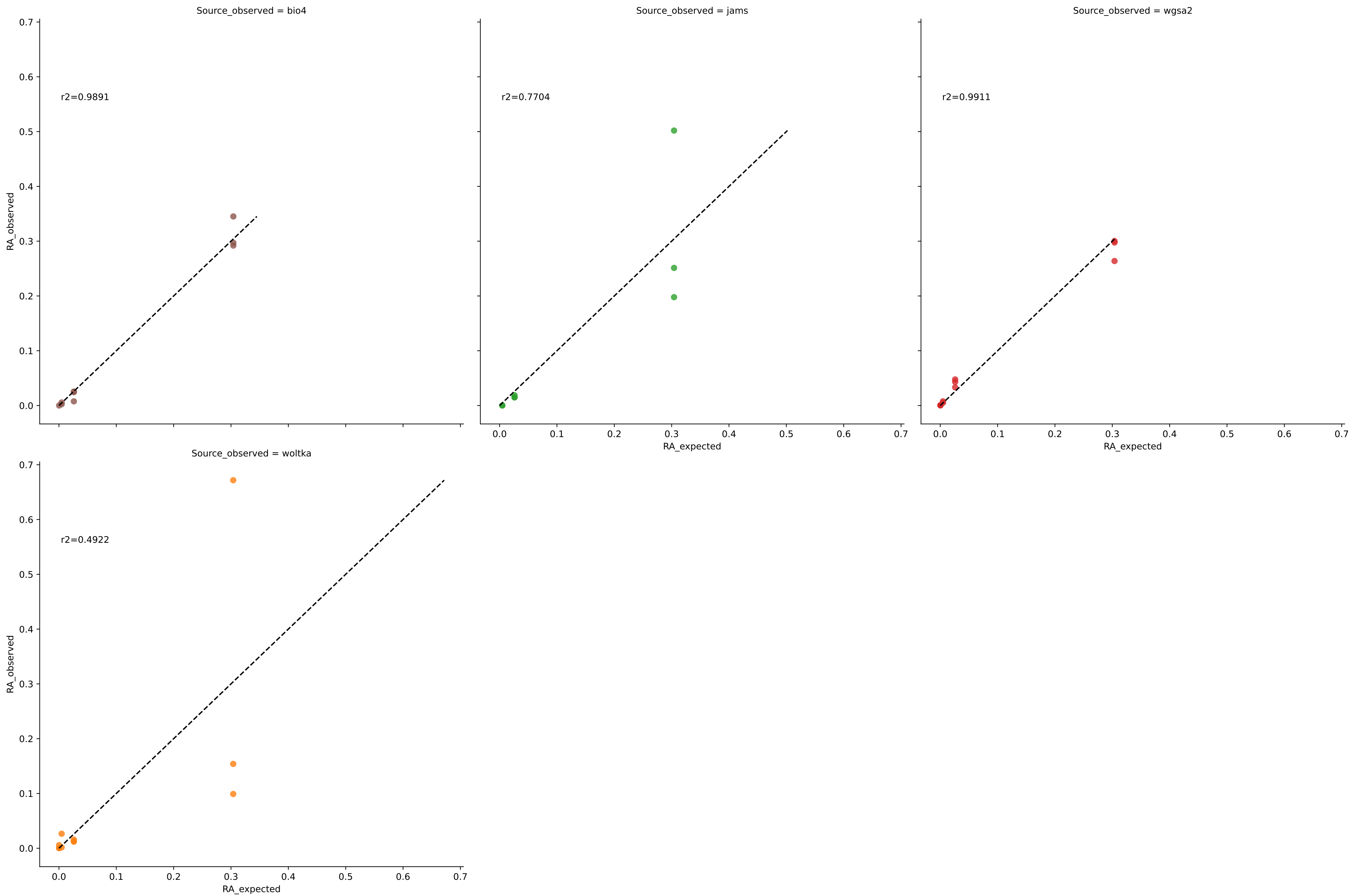


Bivariate Linear Regression for Sample EG in Experiment nist (Genus at filter threshold 0)

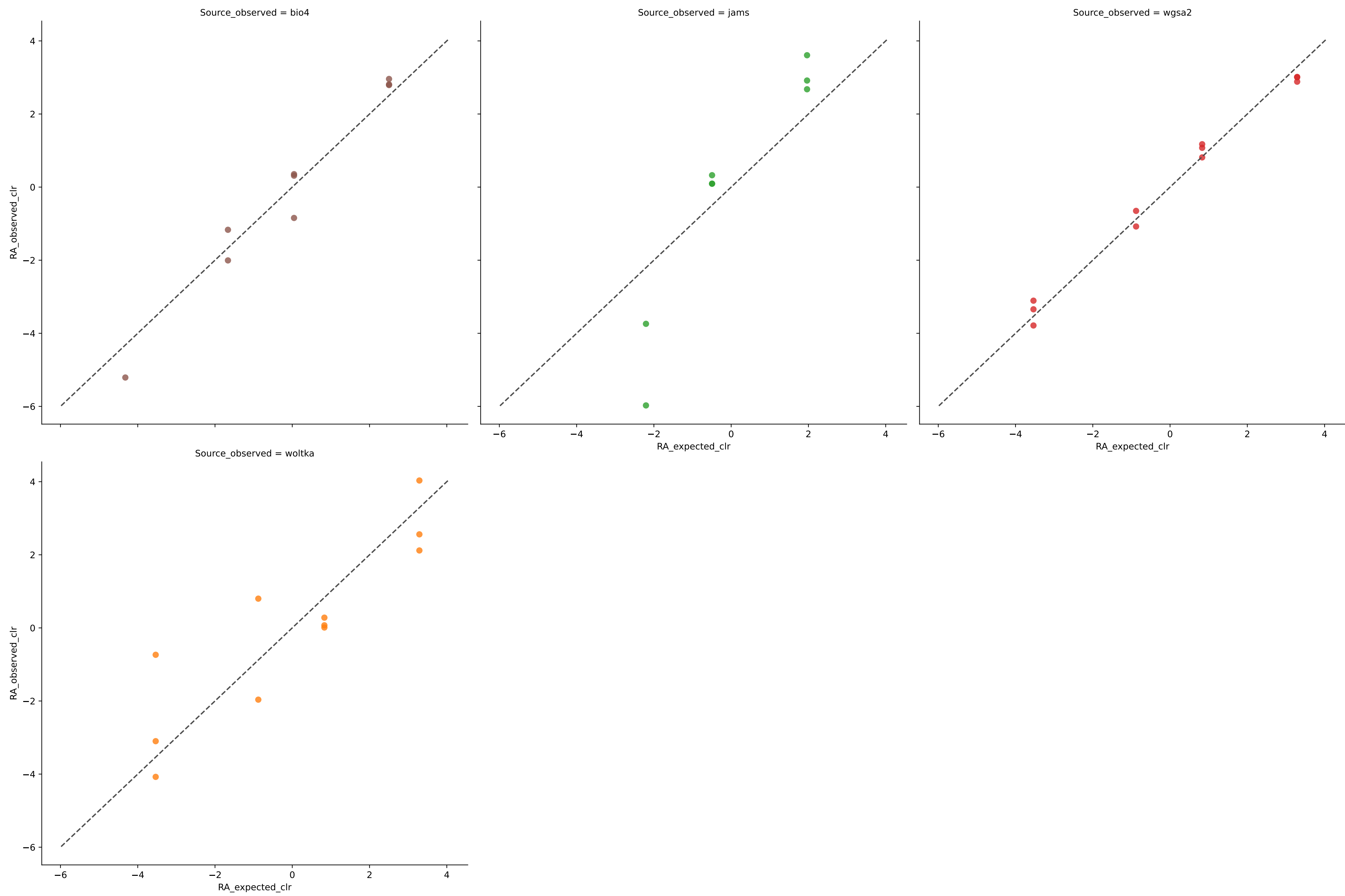


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	12	0.4922	0.0195	1.6584	0.8827	0.0296	100.0000	0.0000
jams	14	0.4230	0.0246	6.7319	0.8279	0.0398	100.0000	0.0000
wgsa2	14	0.5598	0.0202	3.8550	0.8586	0.0276	100.0000	0.0000
woltka	14	0.1669	0.0549	3.6377	0.6157	0.0752	100.0000	0.0000

Bivariate Linear Regression for Sample MIX-A in Experiment nist (Genus at filter threshold 0)

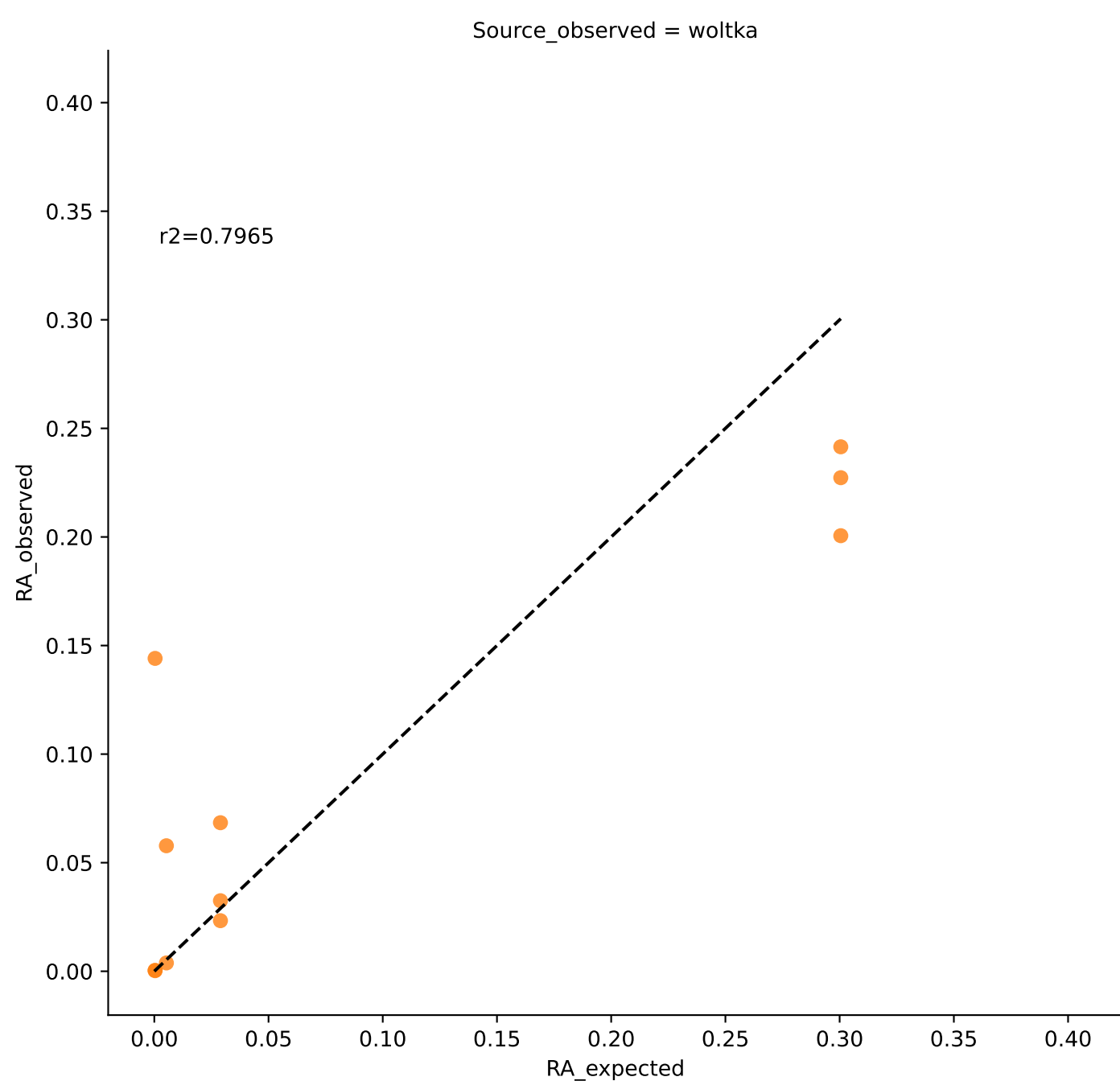
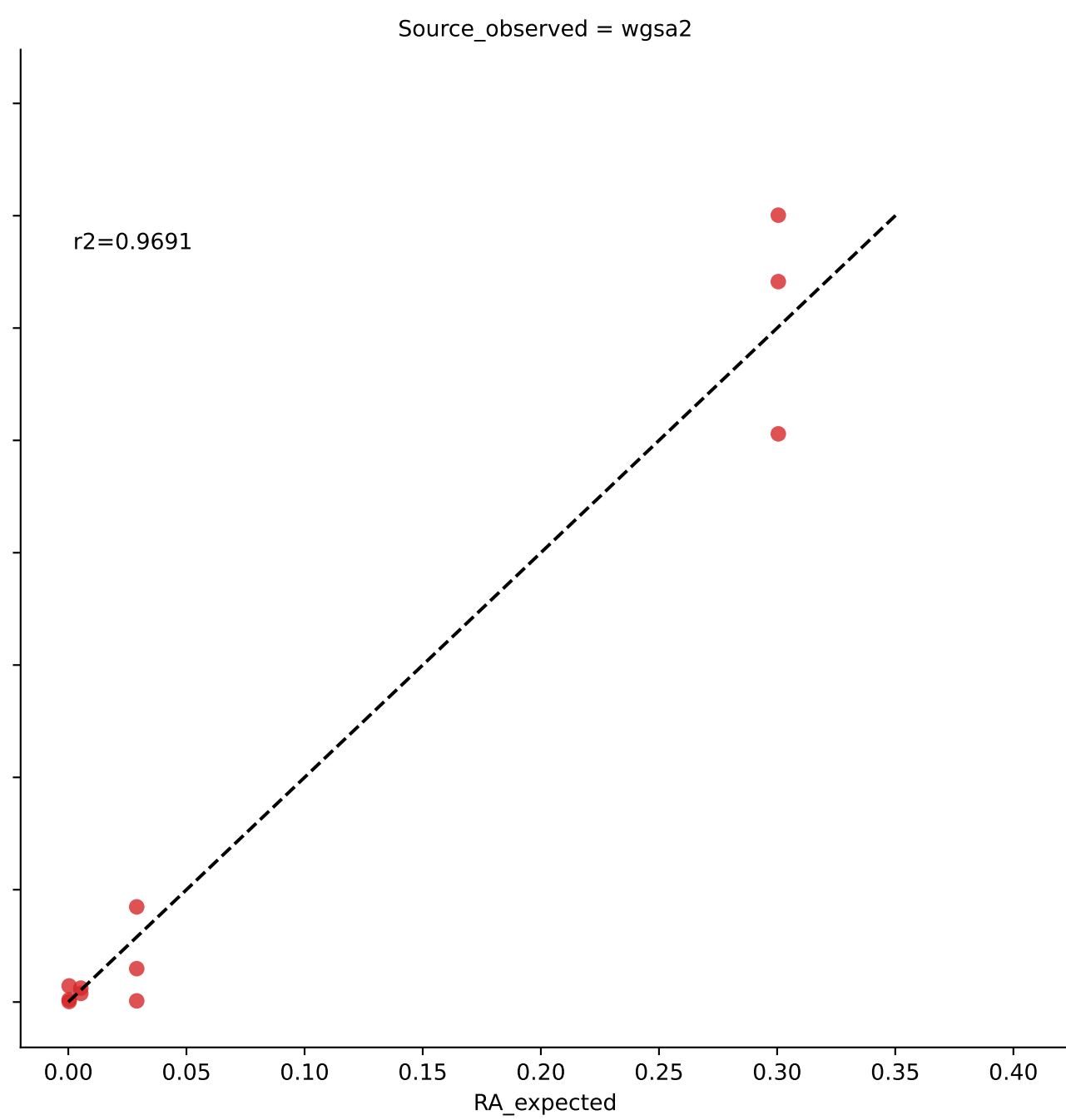
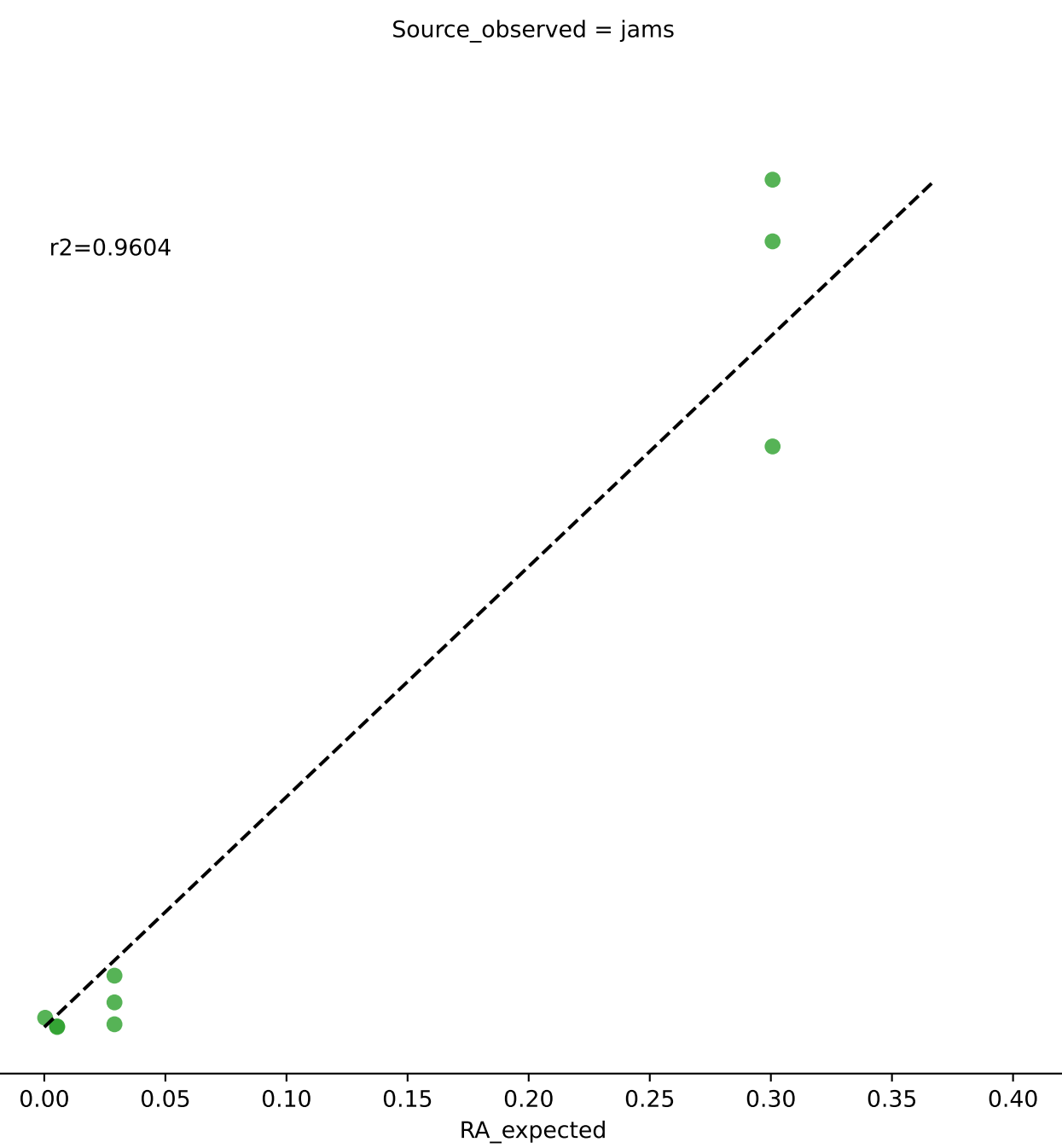
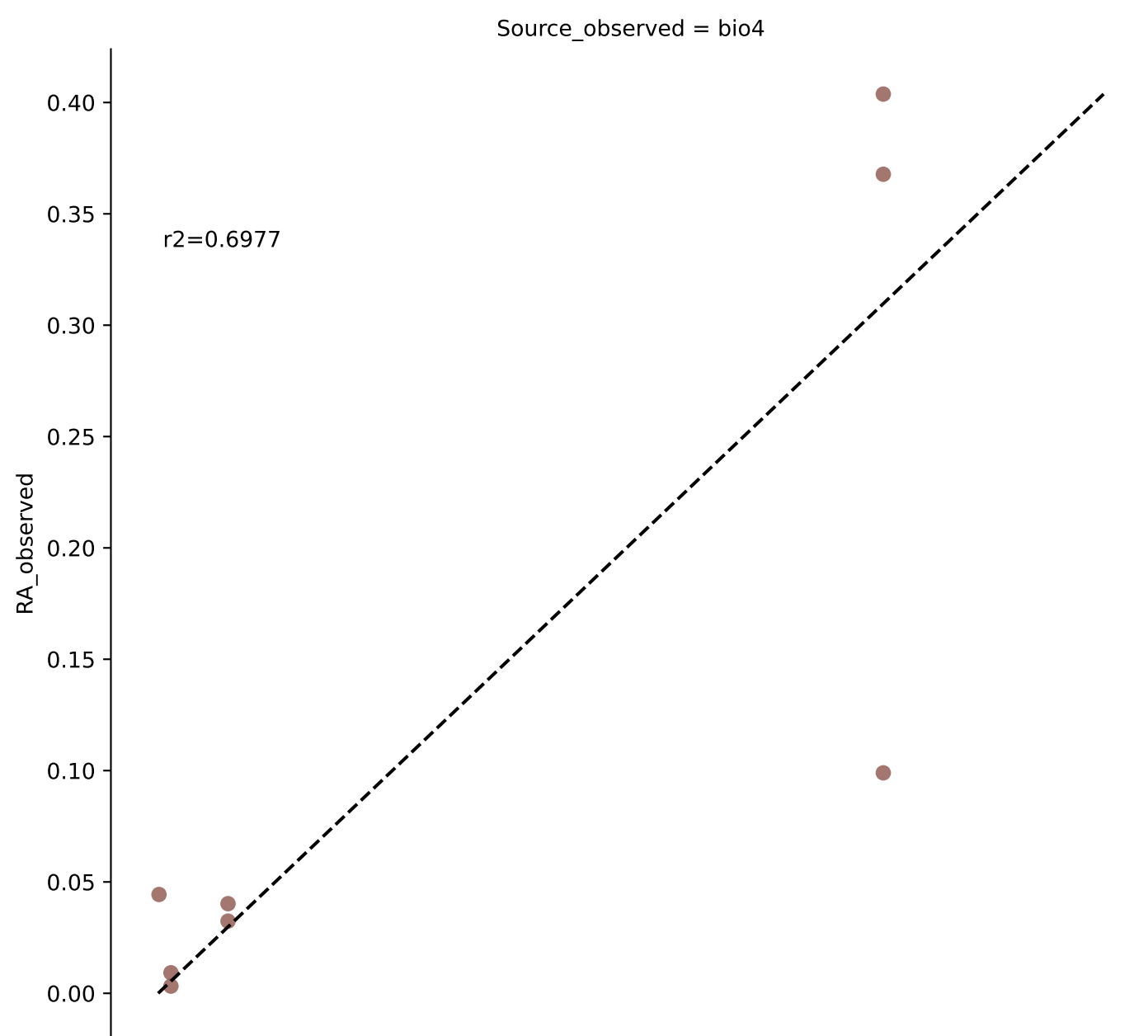


Bivariate Linear Regression for Sample MIX-A in Experiment nist (Genus at filter threshold 0)

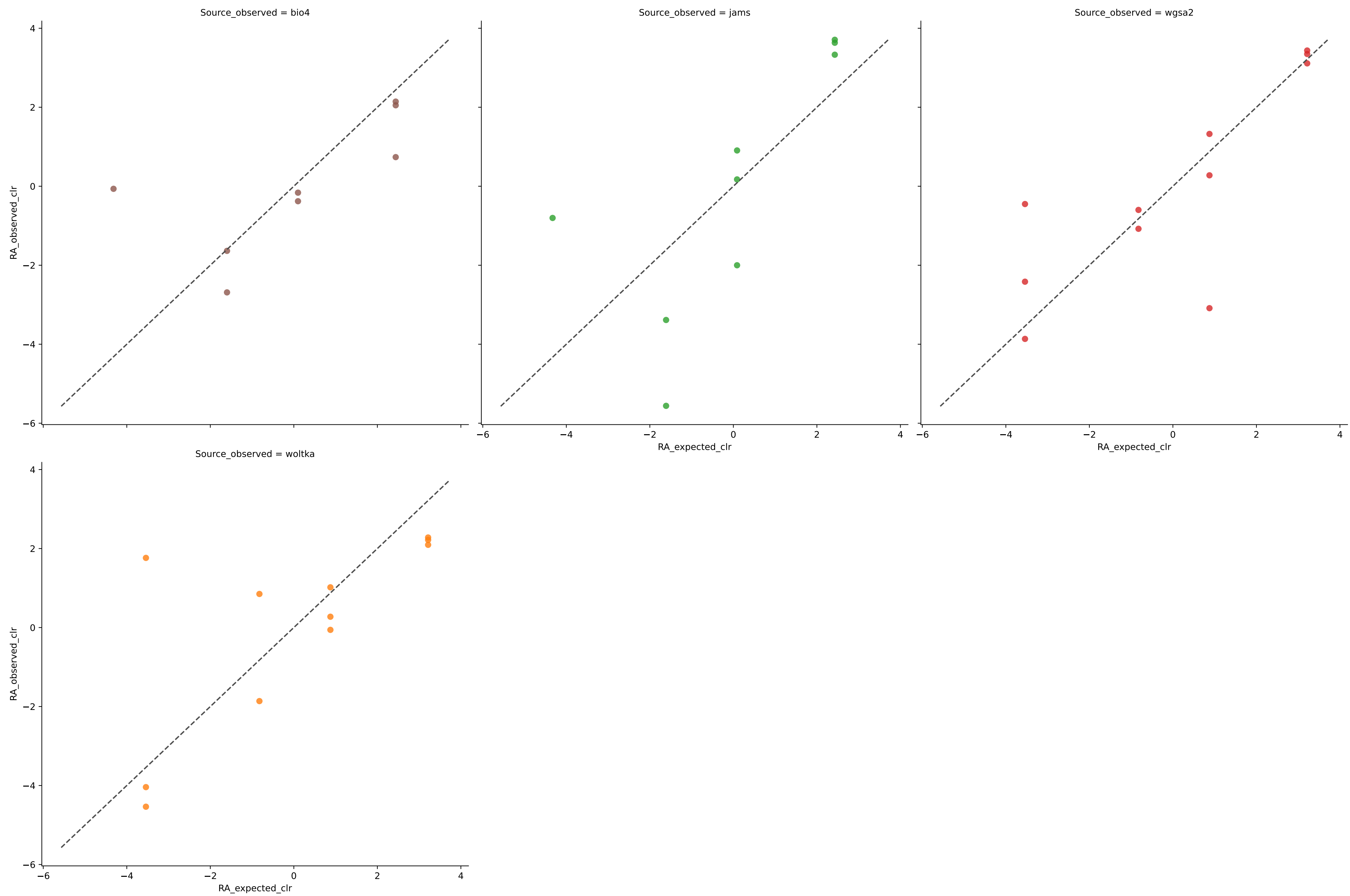


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	9	0.9891	0.0093	1.5789	0.9581	0.0157	100.0000	0.0000
jams	8	0.7704	0.0494	4.6943	0.8023	0.0818	100.0000	0.0000
wgsa2	11	0.9911	0.0091	0.9344	0.9501	0.0150	100.0000	0.0000
woltka	11	0.4922	0.0719	4.0410	0.6048	0.1351	100.0000	0.0000

Bivariate Linear Regression for Sample MIX-B in Experiment nist (Genus at filter threshold 0)

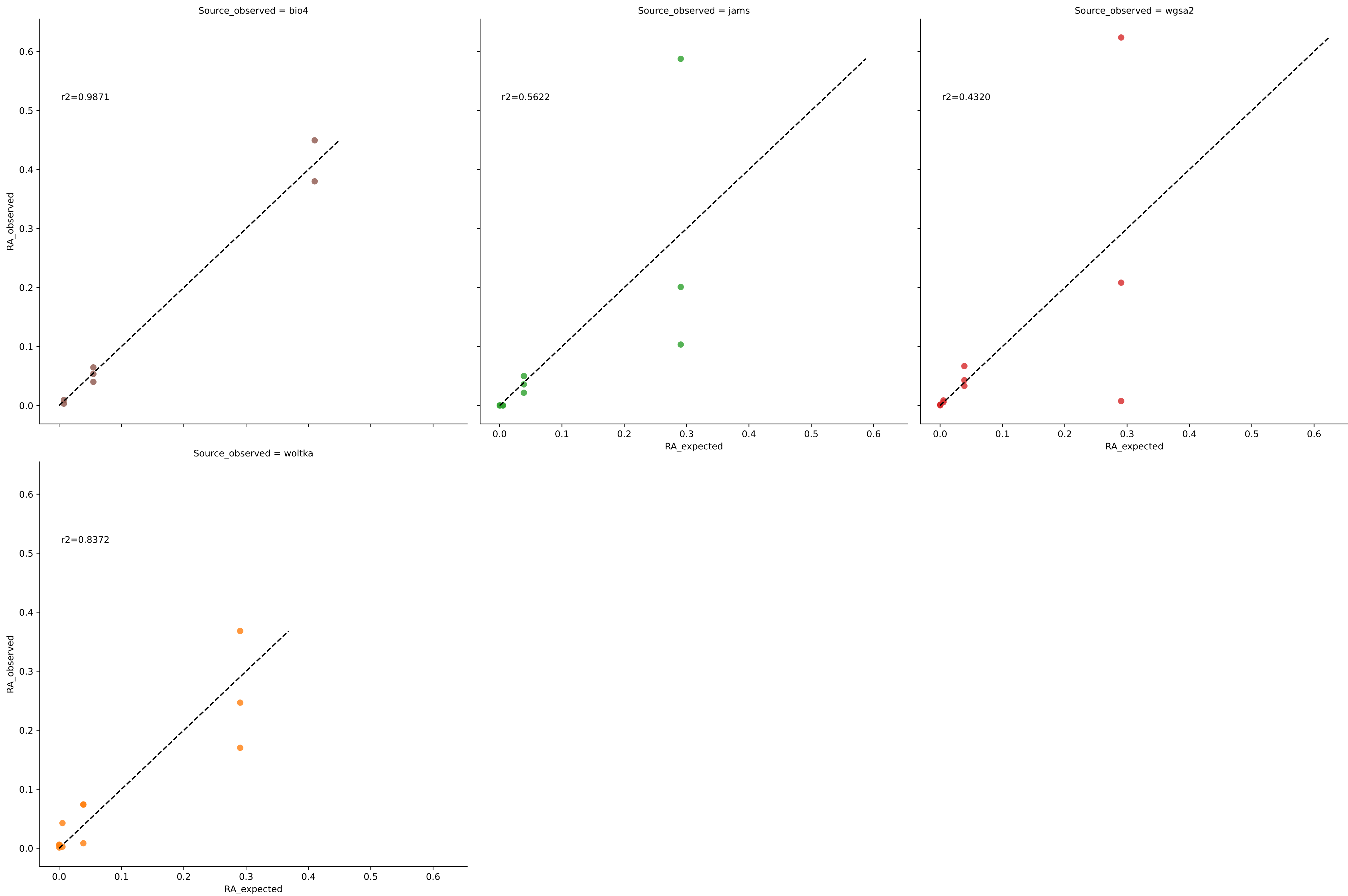


Bivariate Linear Regression for Sample MIX-B in Experiment nist (Genus at filter threshold 0)

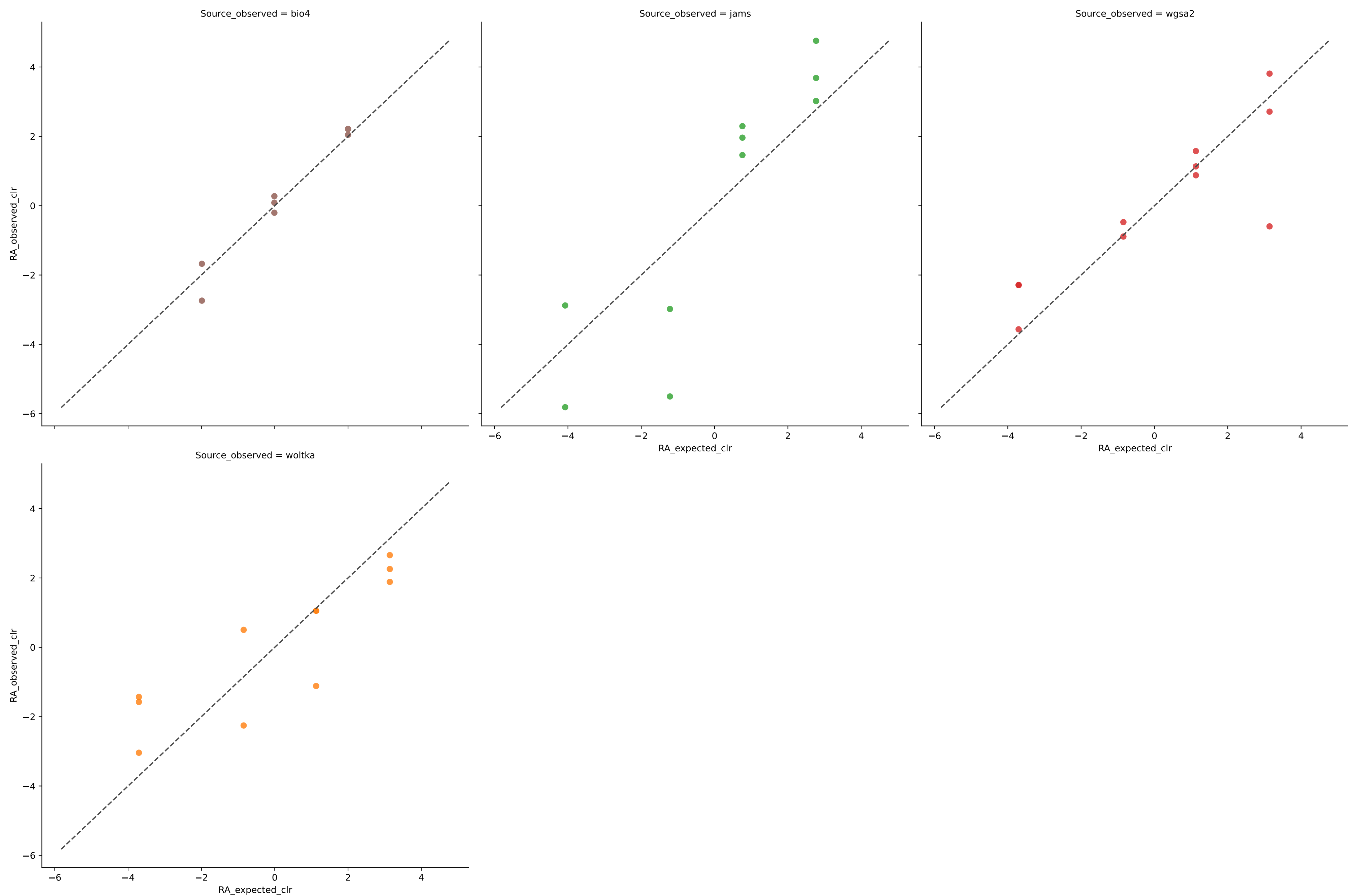


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	8	0.6977	0.0532	4.7646	0.7871	0.0857	100.0000	0.0000
jams	9	0.9604	0.0248	6.3339	0.8885	0.0329	100.0000	0.0000
wgsa2	11	0.9691	0.0167	5.2327	0.9083	0.0241	100.0000	0.0000
woltka	11	0.7965	0.0435	6.1353	0.7607	0.0631	100.0000	0.0000

Bivariate Linear Regression for Sample MIX-C in Experiment nist (Genus at filter threshold 0)

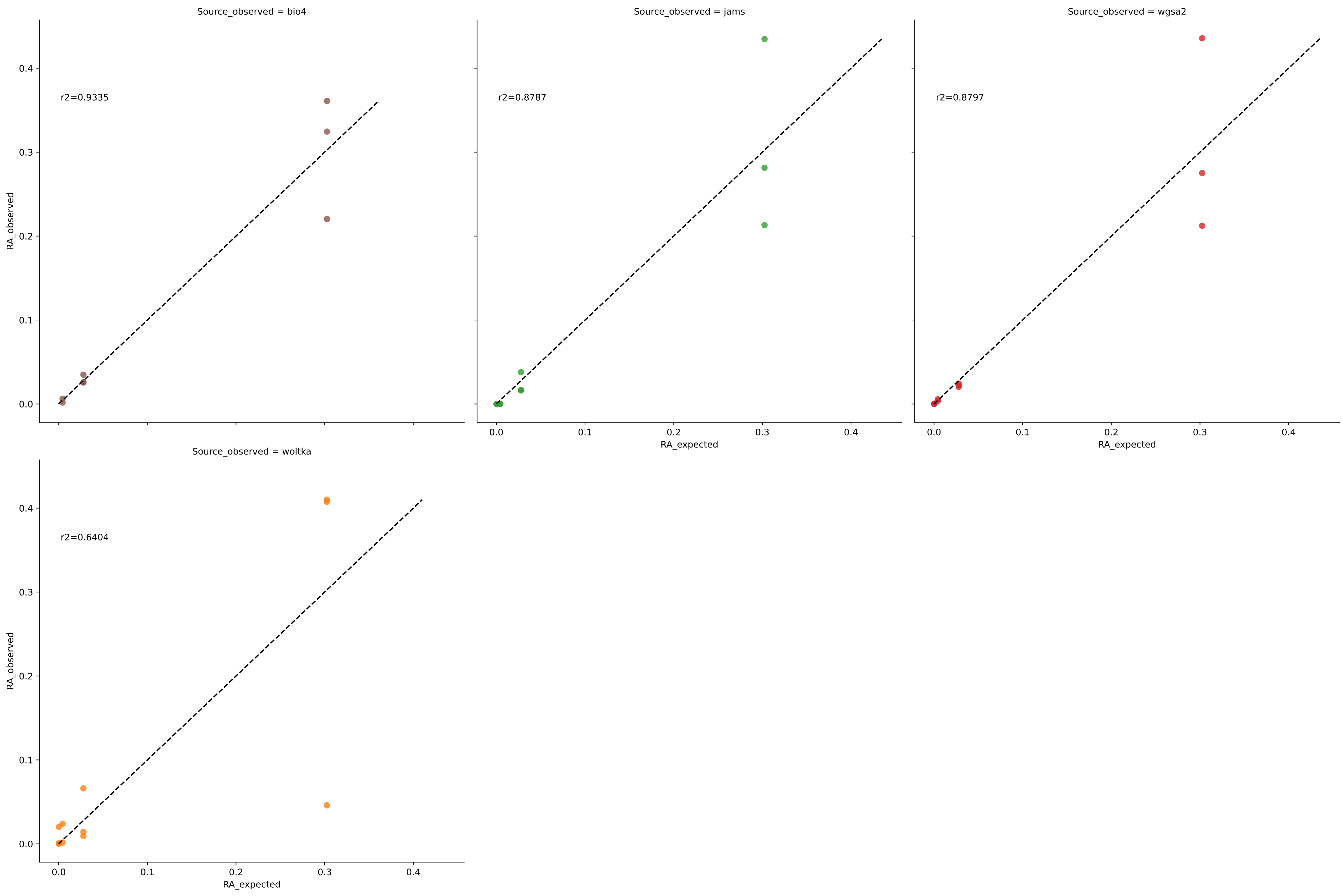


Bivariate Linear Regression for Sample MIX-C in Experiment nist (Genus at filter threshold 0)

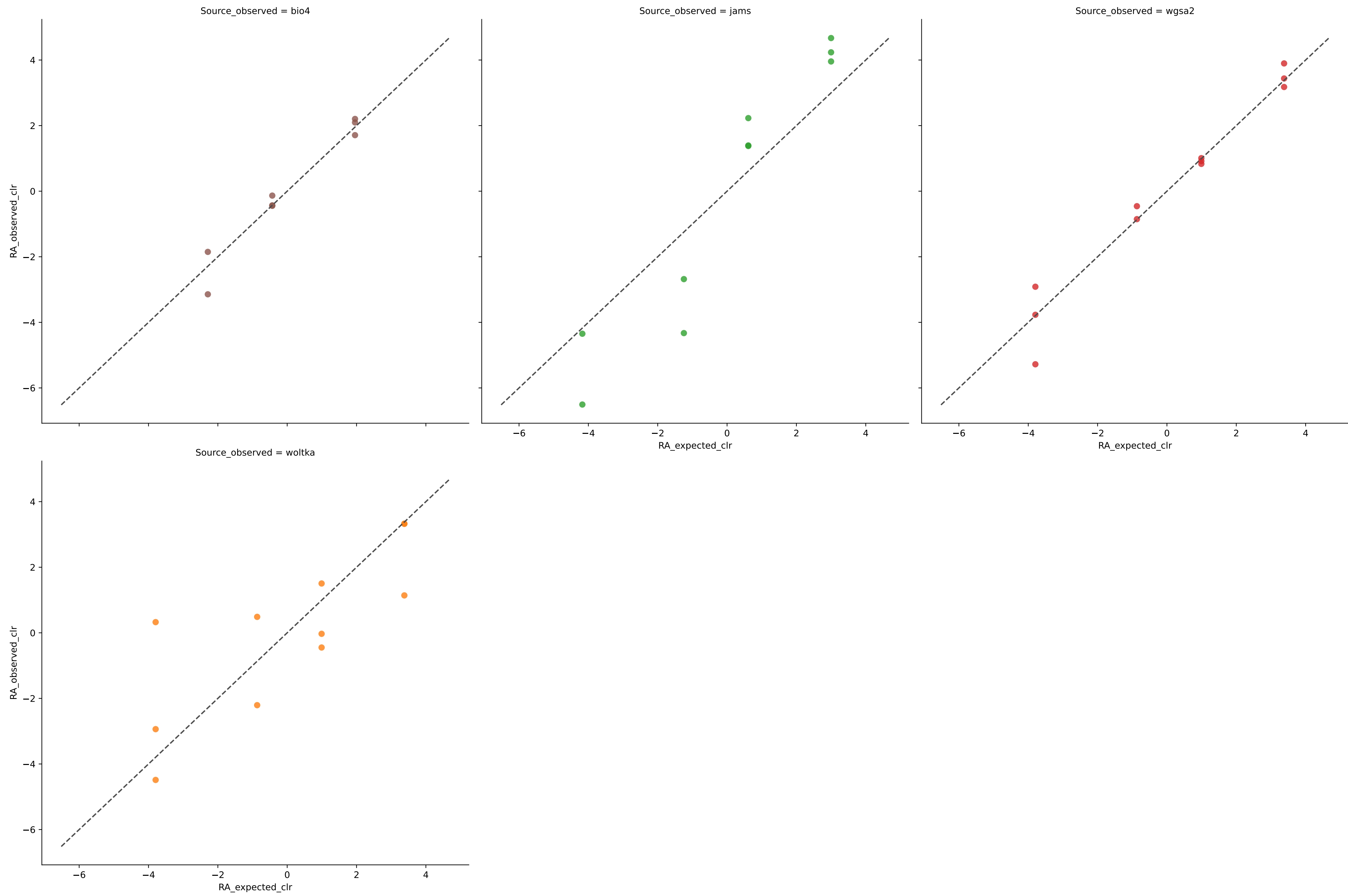


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	7	0.9871	0.0145	0.9149	0.9493	0.0200	100.0000	0.0000
jams	10	0.5622	0.0616	5.9201	0.6920	0.1148	100.0000	0.0000
wgsa2	11	0.4320	0.0674	4.3618	0.6292	0.1344	100.0000	0.0000
woltka	11	0.8372	0.0358	4.6458	0.8029	0.0498	100.0000	0.0000

Bivariate Linear Regression for Sample MIX-D in Experiment nist (Genus at filter threshold 0)

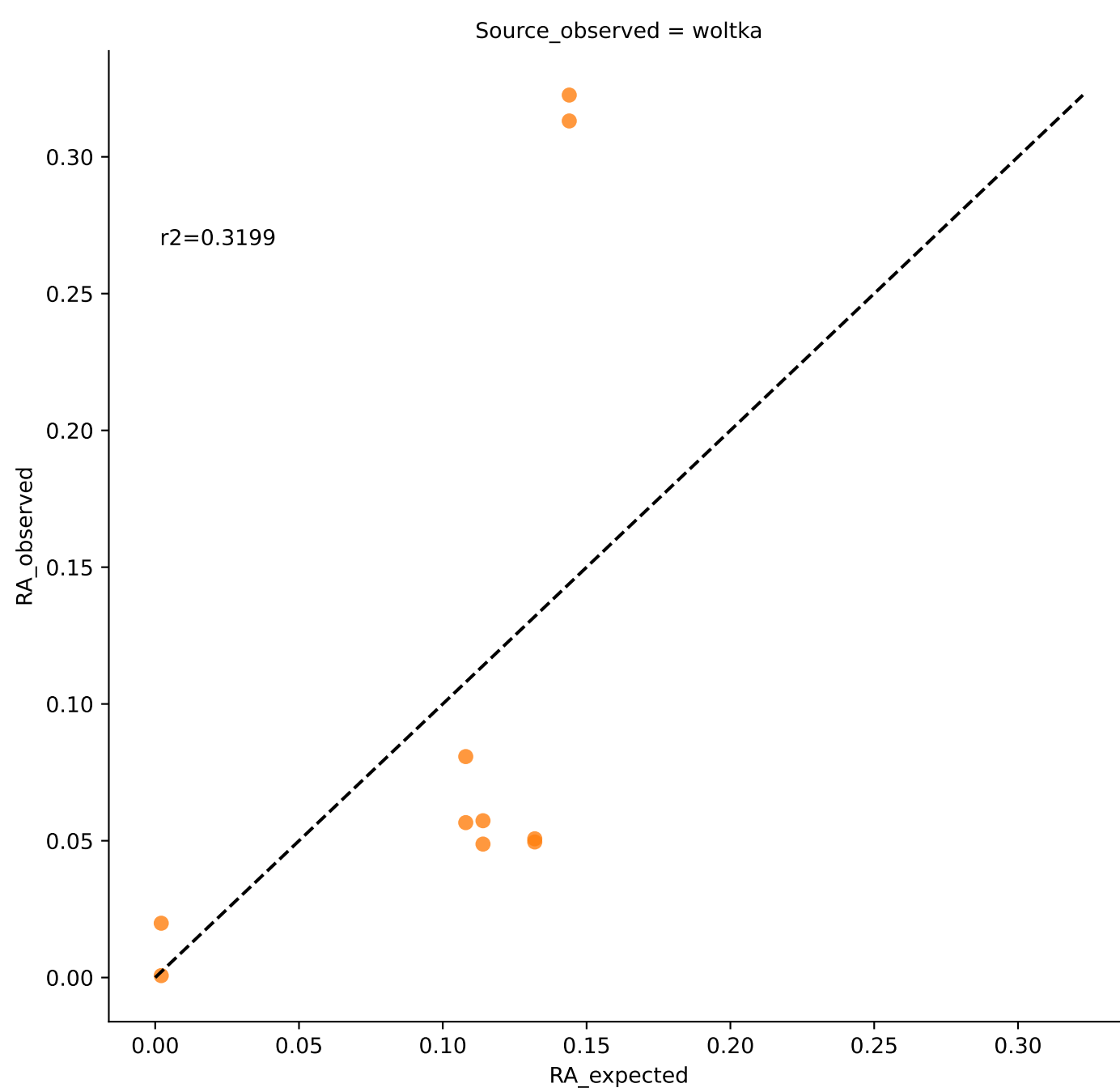
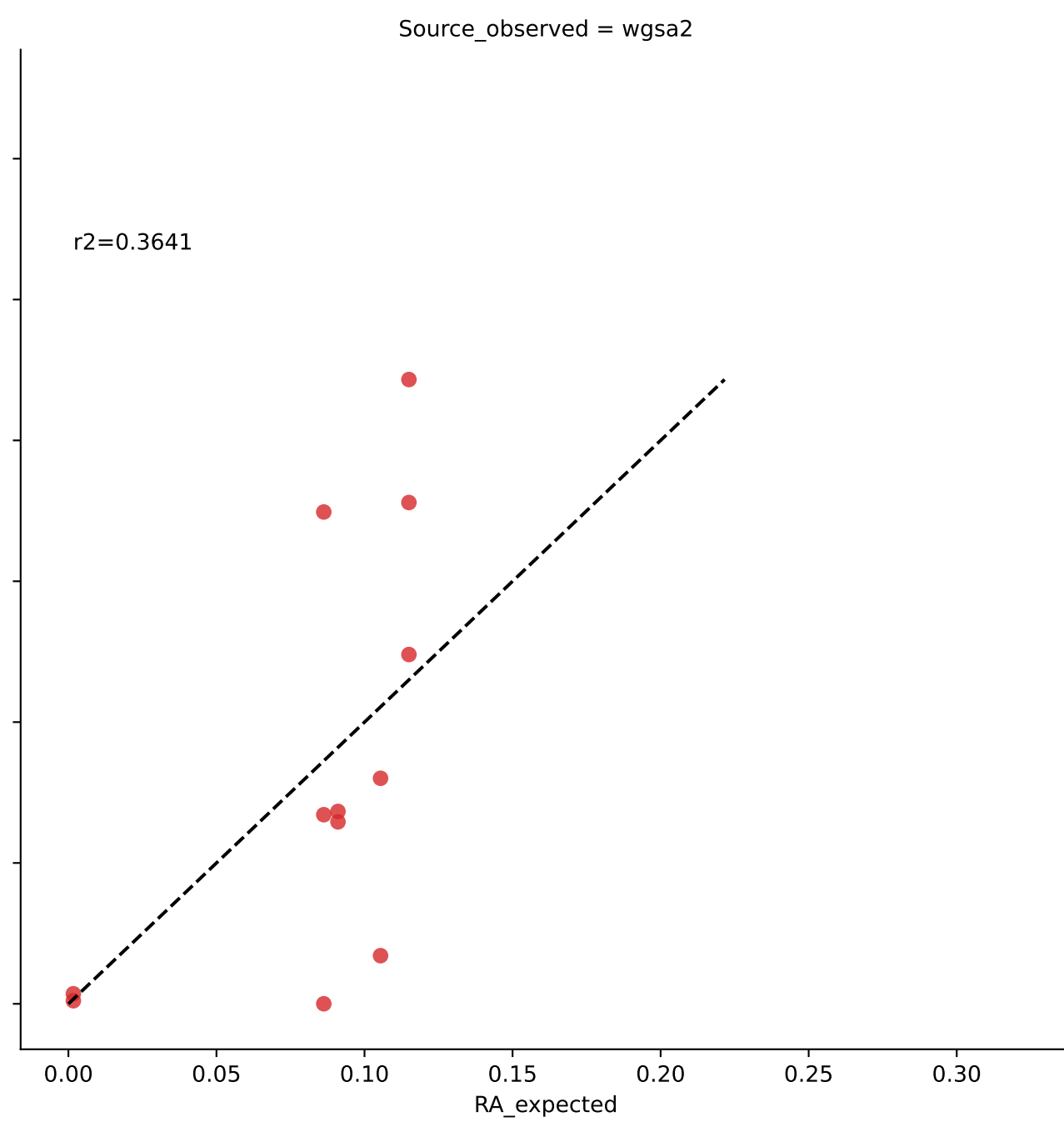
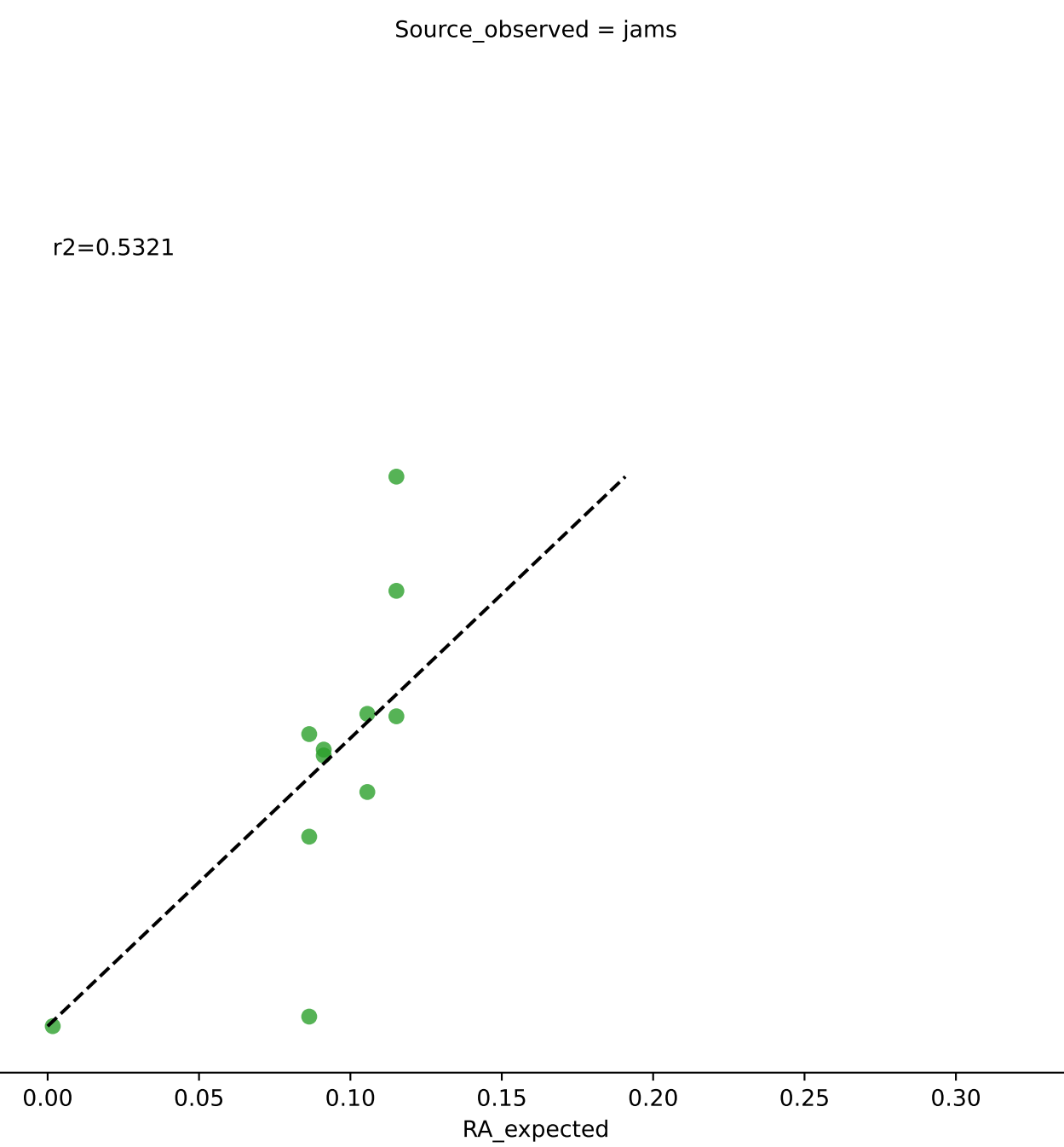
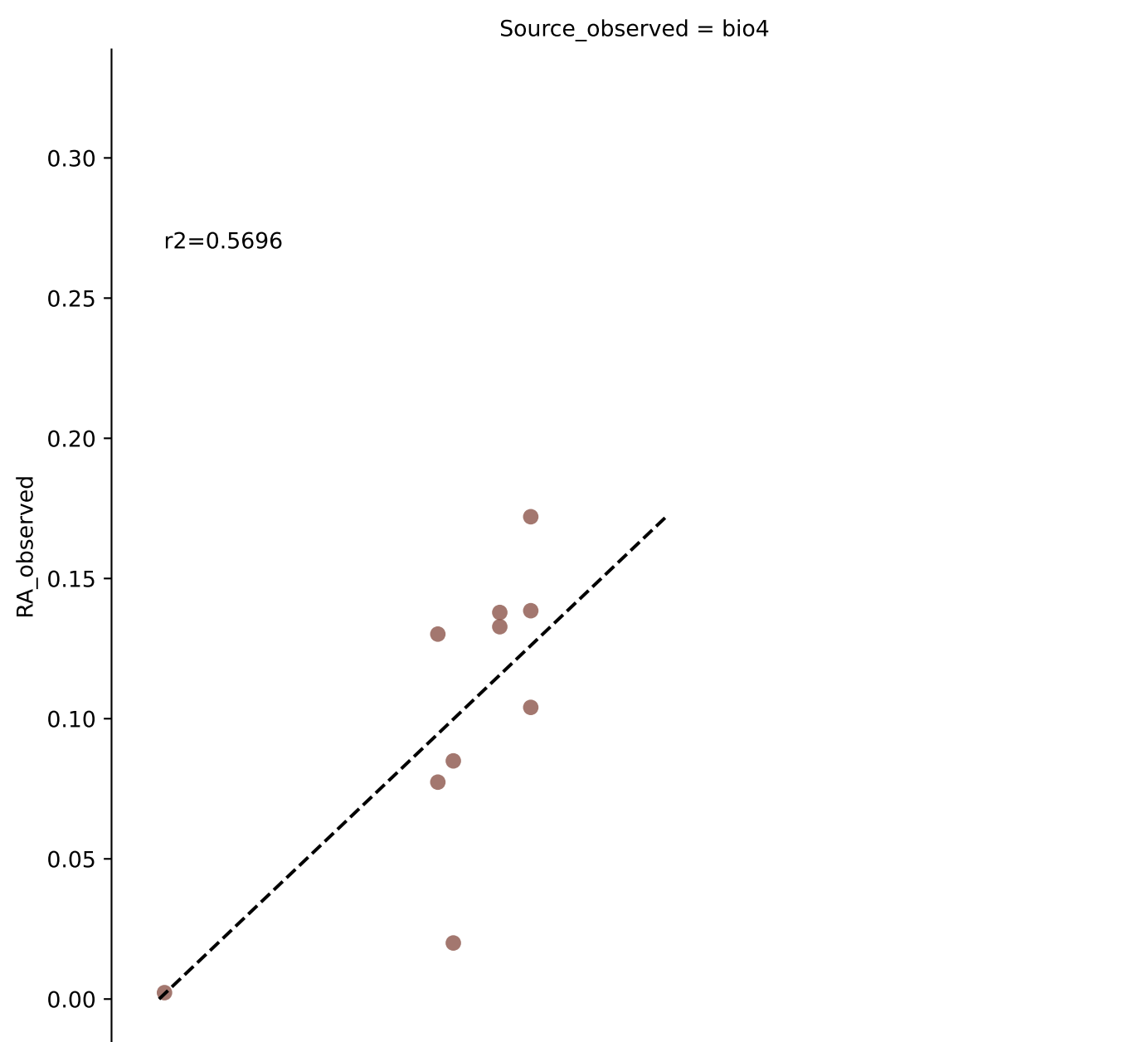


Bivariate Linear Regression for Sample MIX-D in Experiment nist (Genus at filter threshold 0)

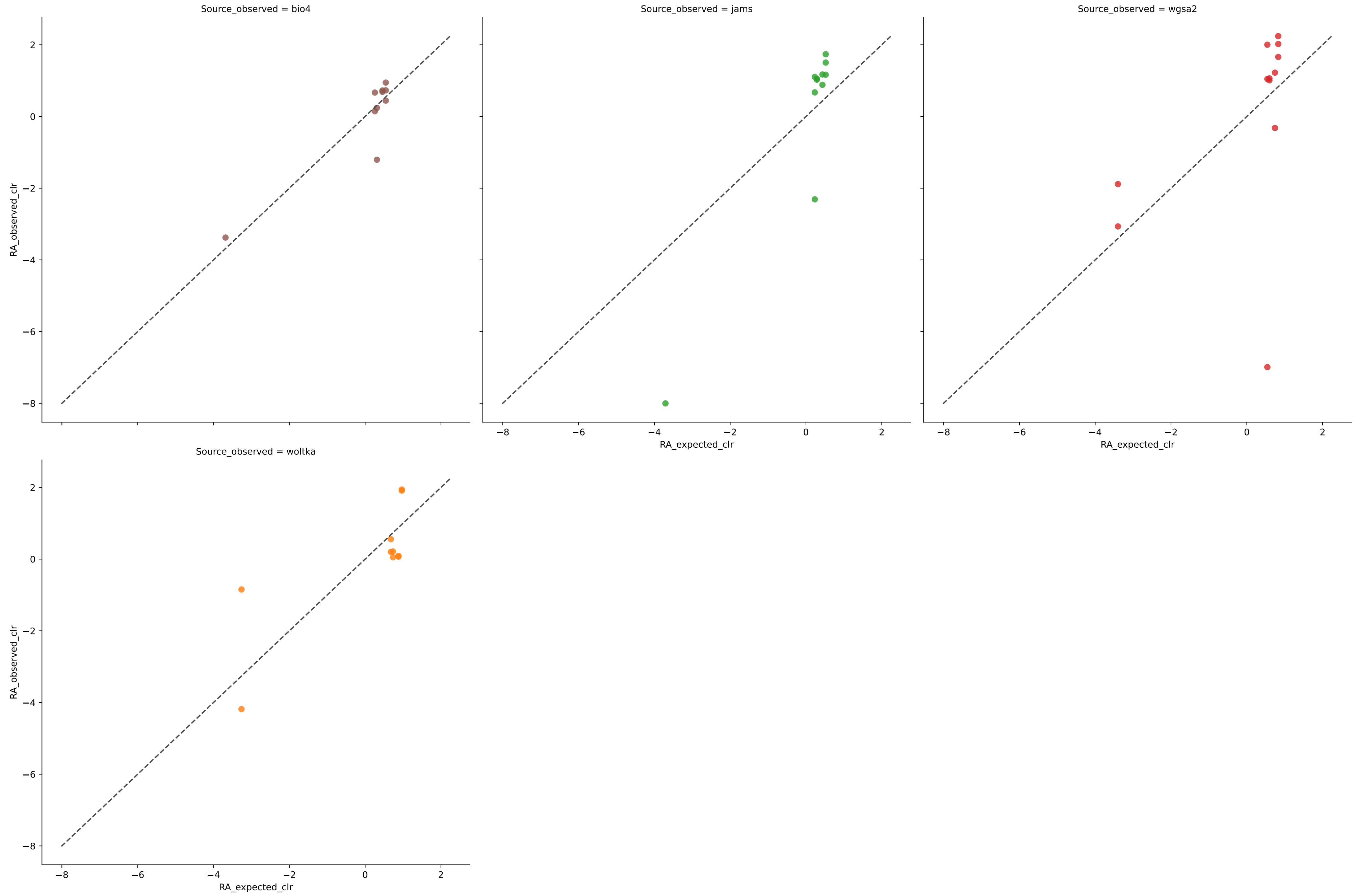


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	8	0.9335	0.0223	1.0739	0.9107	0.0367	100.0000	0.0000
jams	10	0.8787	0.0285	5.1062	0.8577	0.0513	100.0000	0.0000
wgsa2	11	0.8797	0.0245	1.8678	0.8652	0.0493	100.0000	0.0000
woltka	11	0.6404	0.0529	5.4967	0.7088	0.0910	100.0000	0.0000

Bivariate Linear Regression for Sample EG in Experiment nist (Species at filter threshold 0)

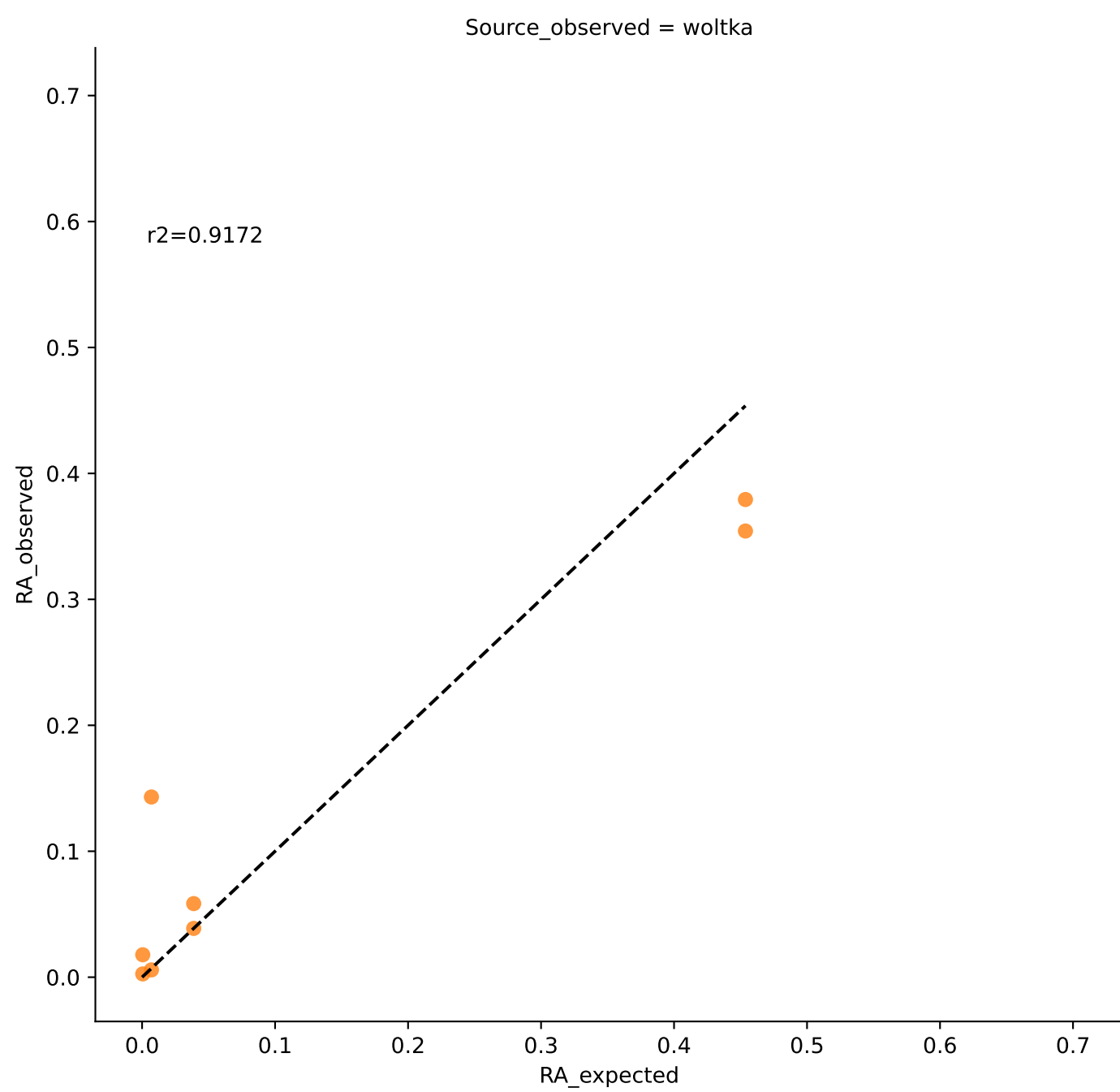
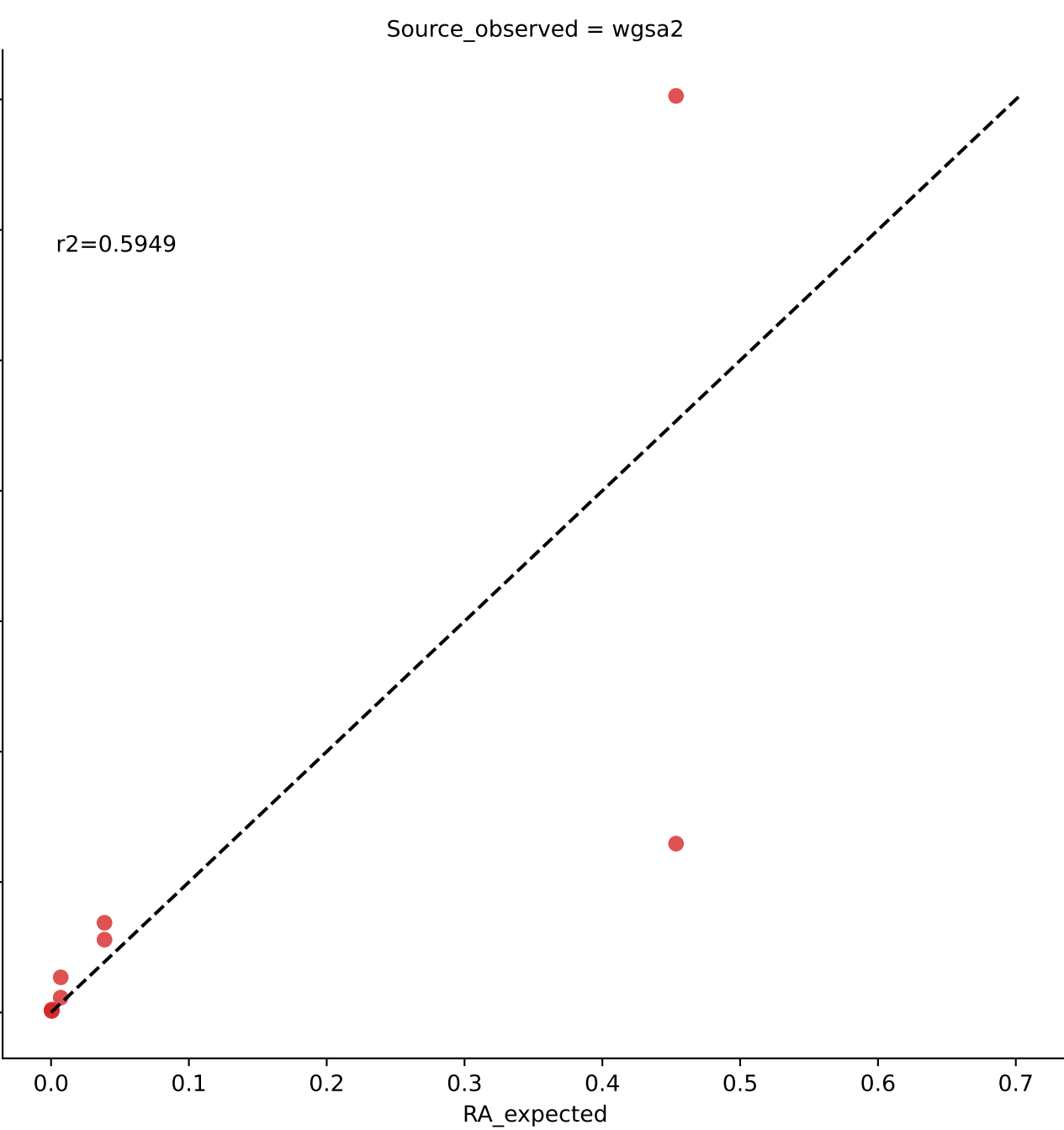
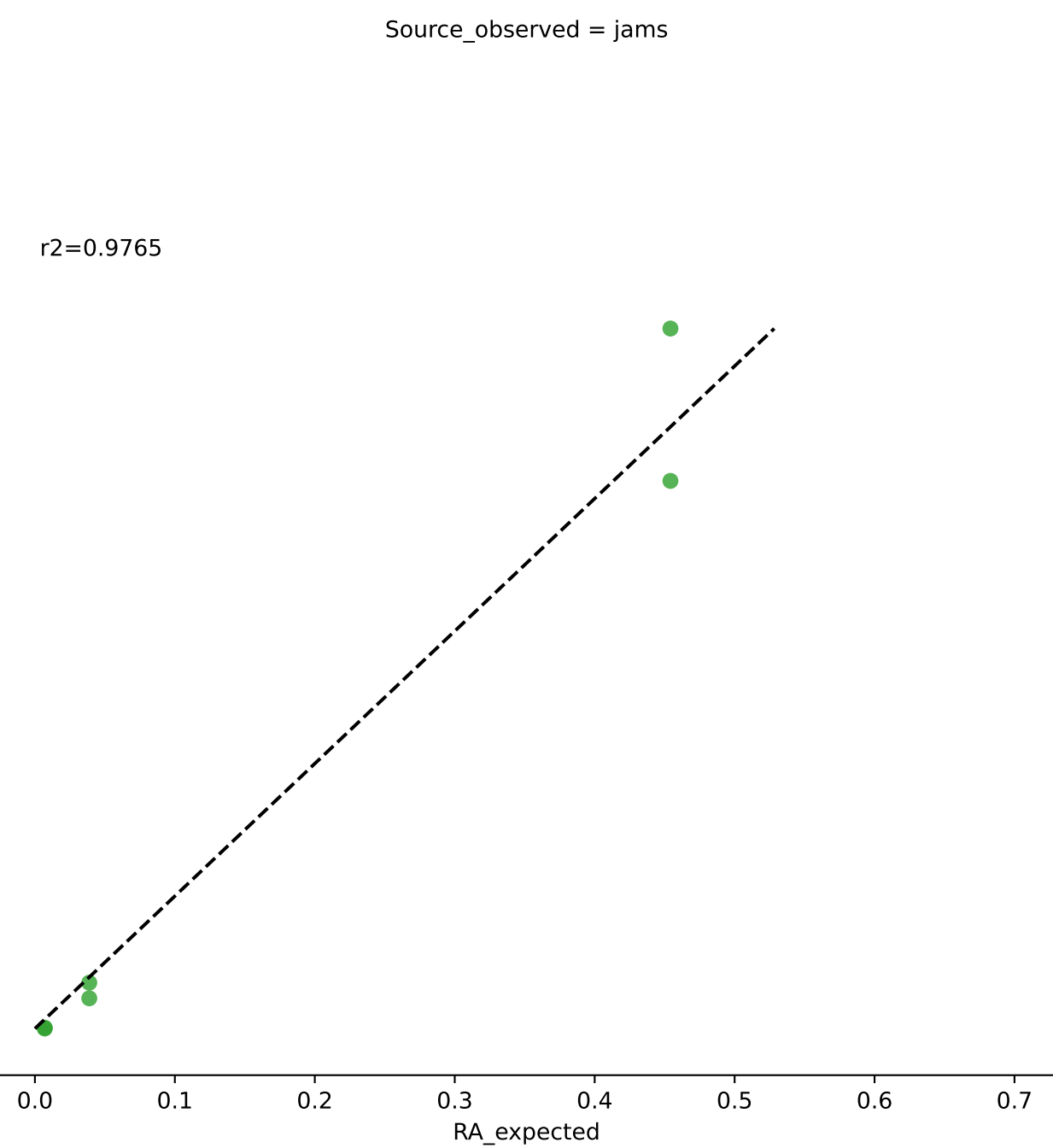
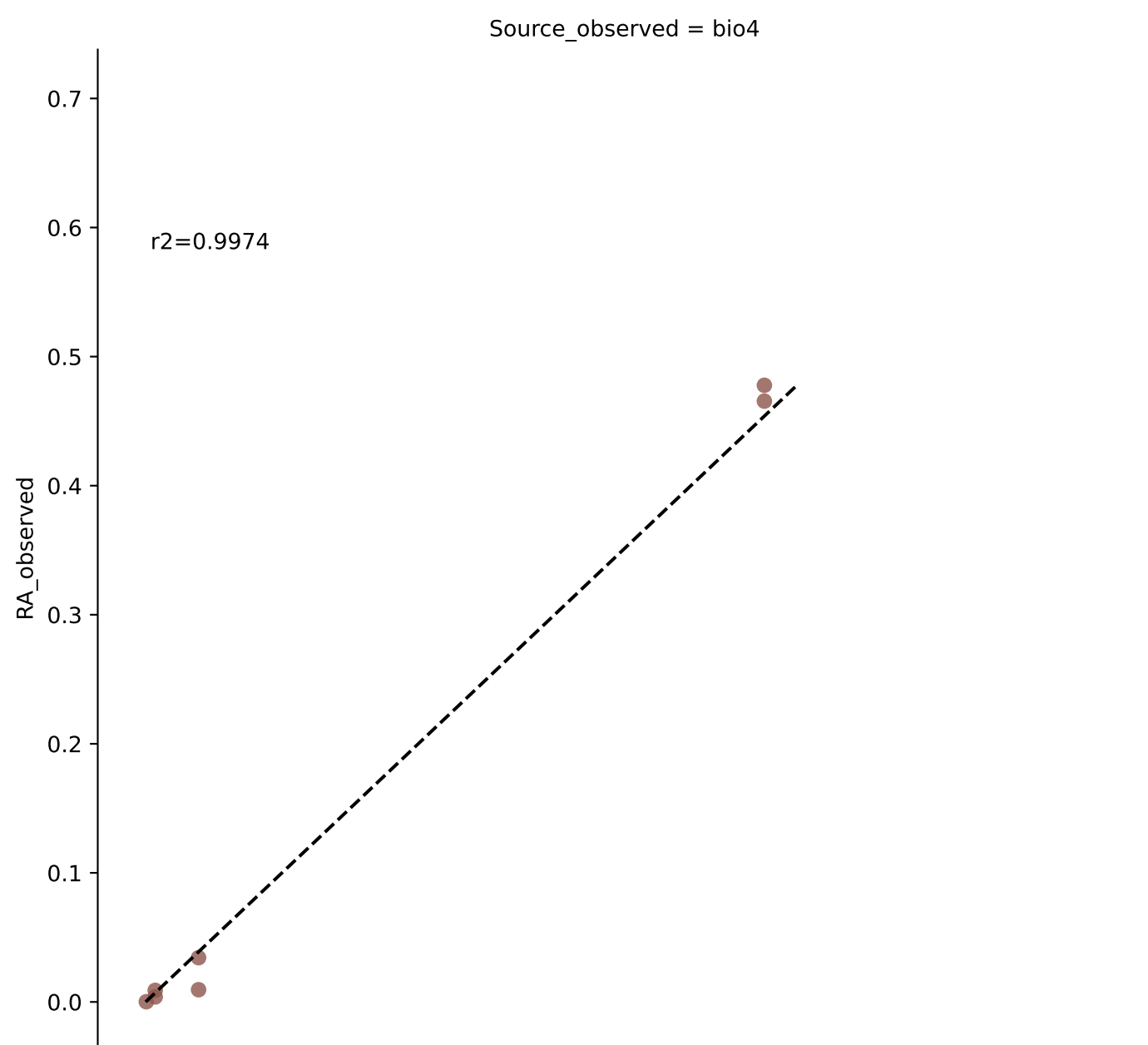


Bivariate Linear Regression for Sample EG in Experiment nist (Species at filter threshold 0)

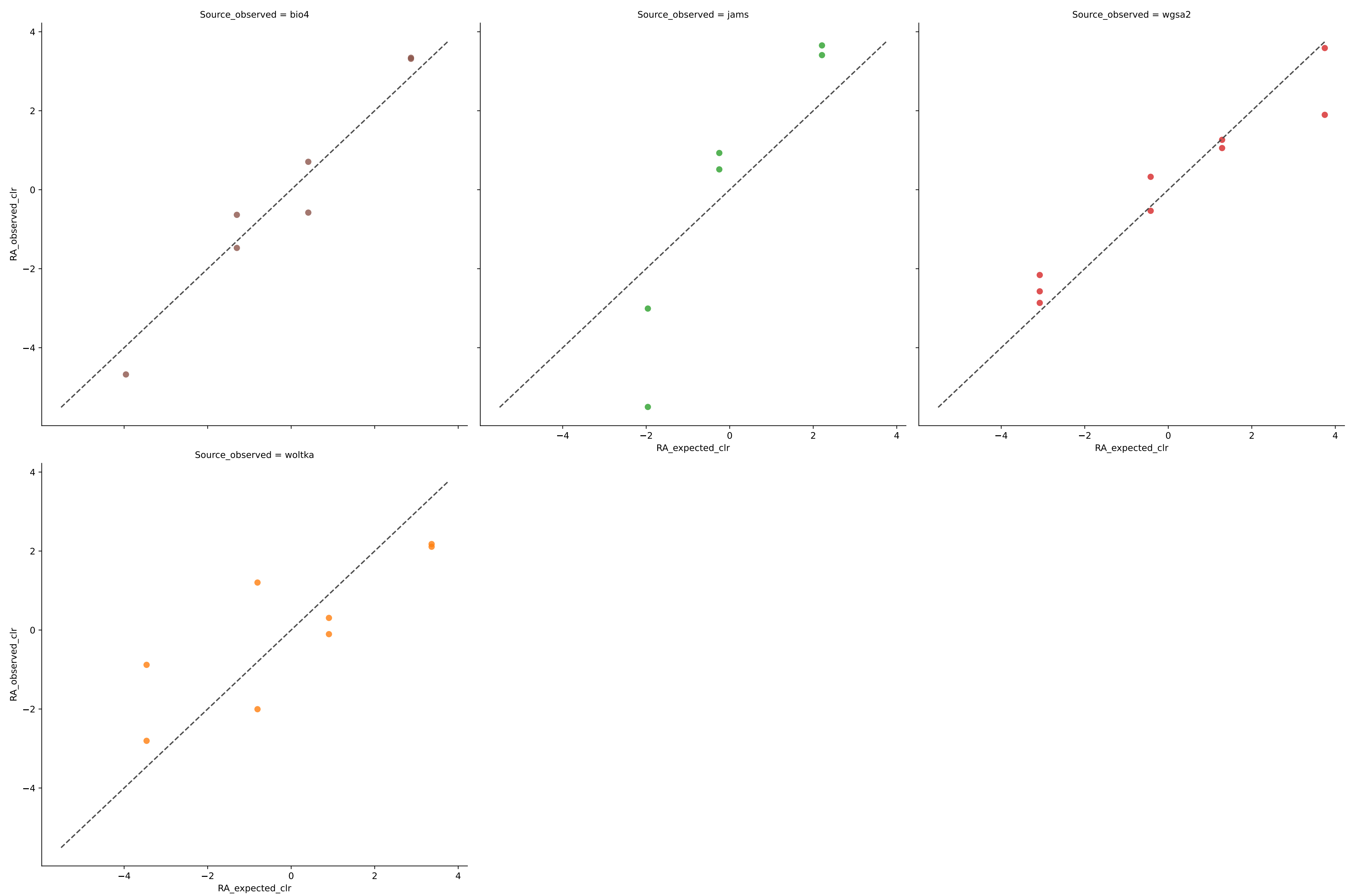


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	10	0.5696	0.0268	1.7058	0.8661	0.0342	100.0000	0.0000
jams	11	0.5321	0.0249	5.5328	0.8628	0.0372	100.0000	0.0000
wgsa2	12	0.3641	0.0448	8.2081	0.7312	0.0582	100.0000	0.0000
woltka	10	0.3199	0.0731	3.2836	0.6345	0.0922	100.0000	0.0000

Bivariate Linear Regression for Sample MIX-A in Experiment nist (Species at filter threshold 0)

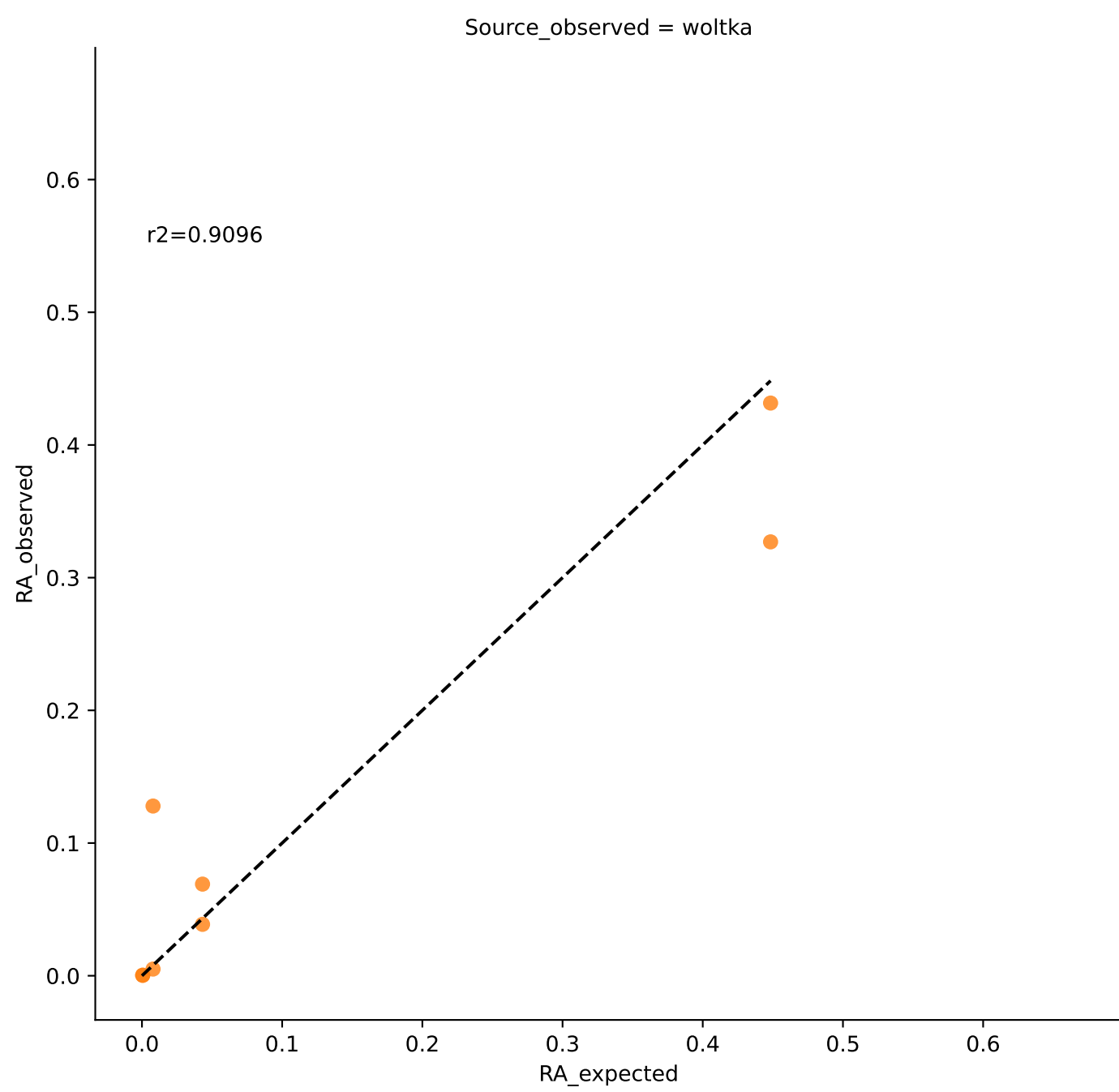
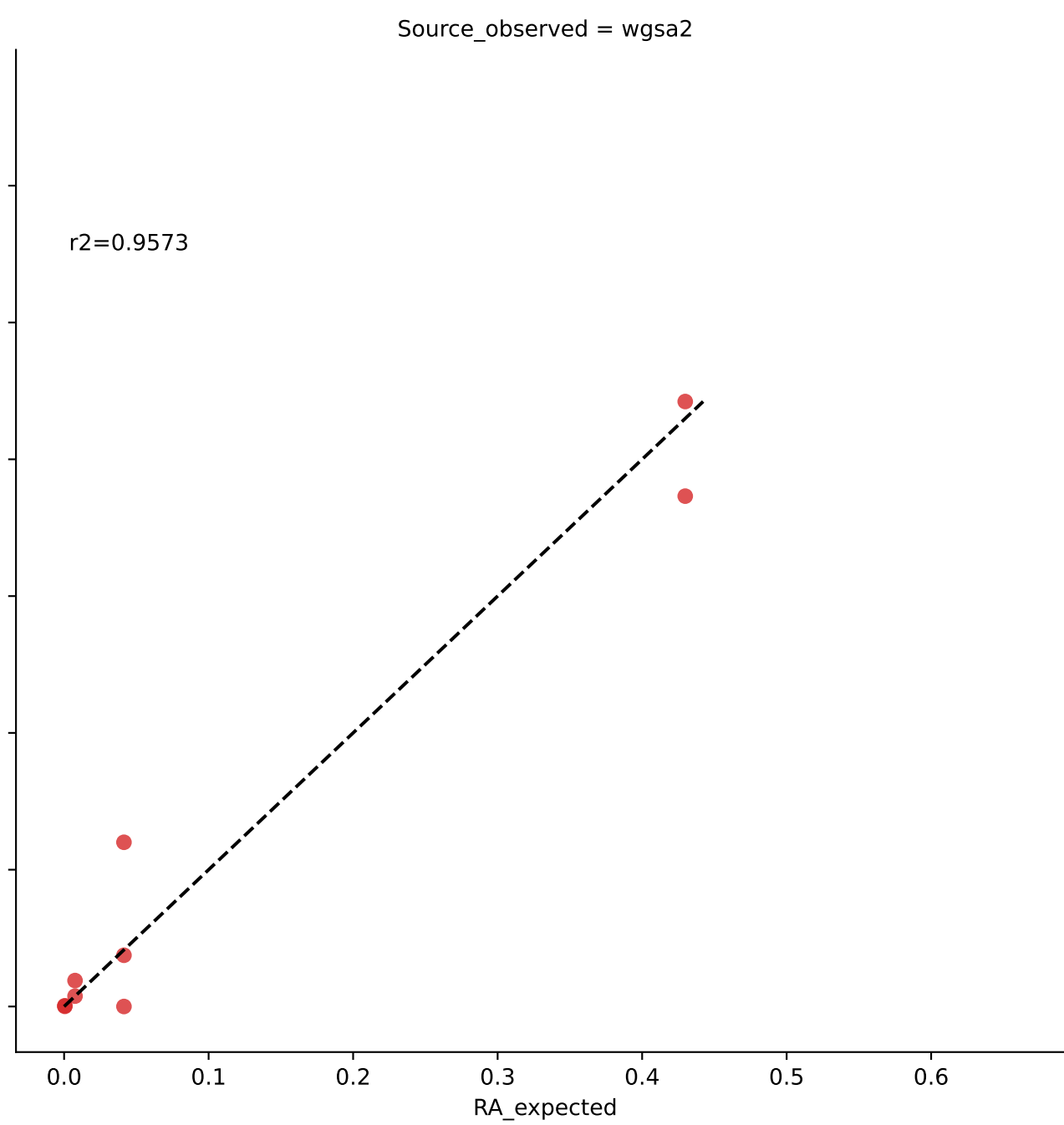
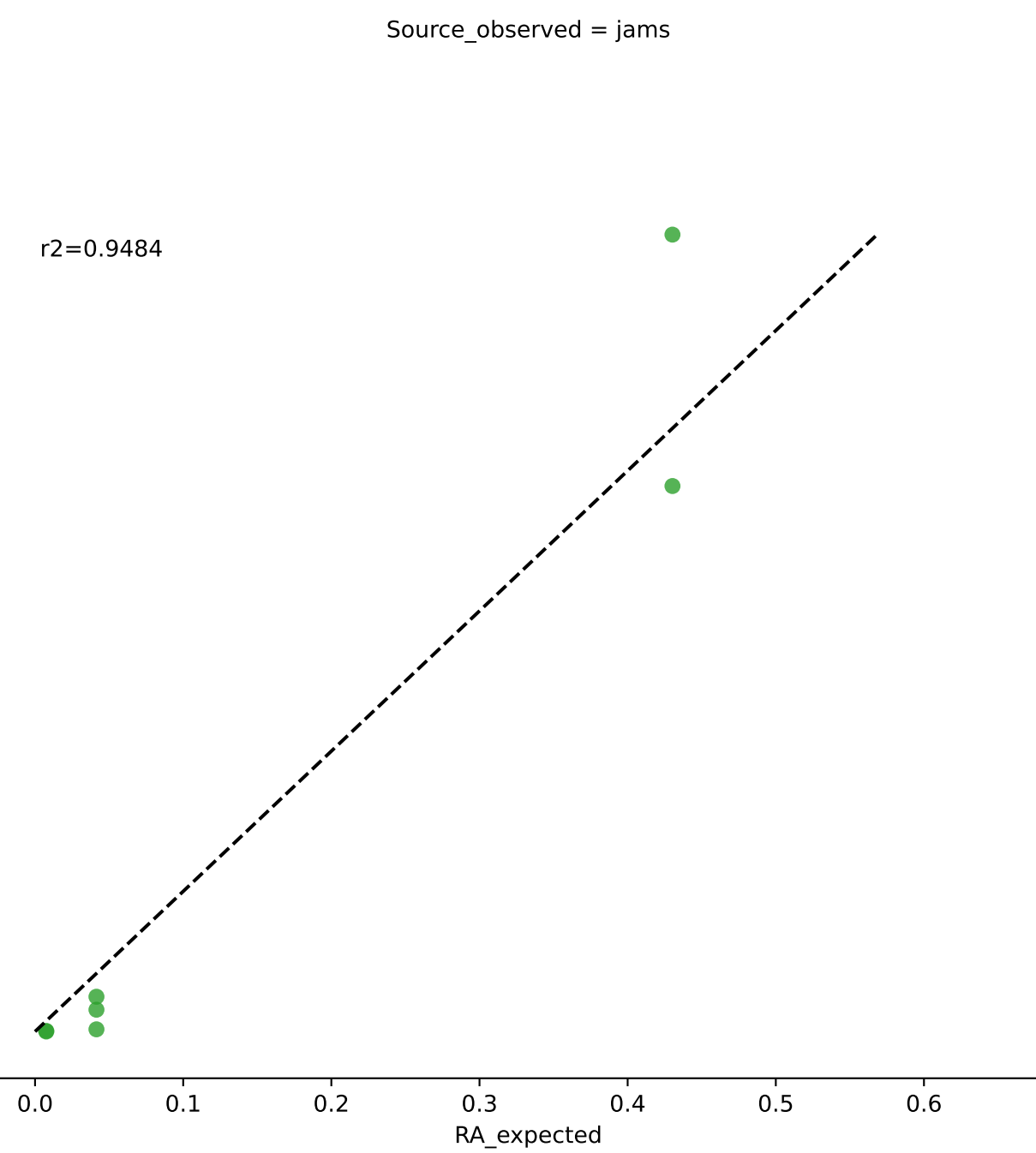
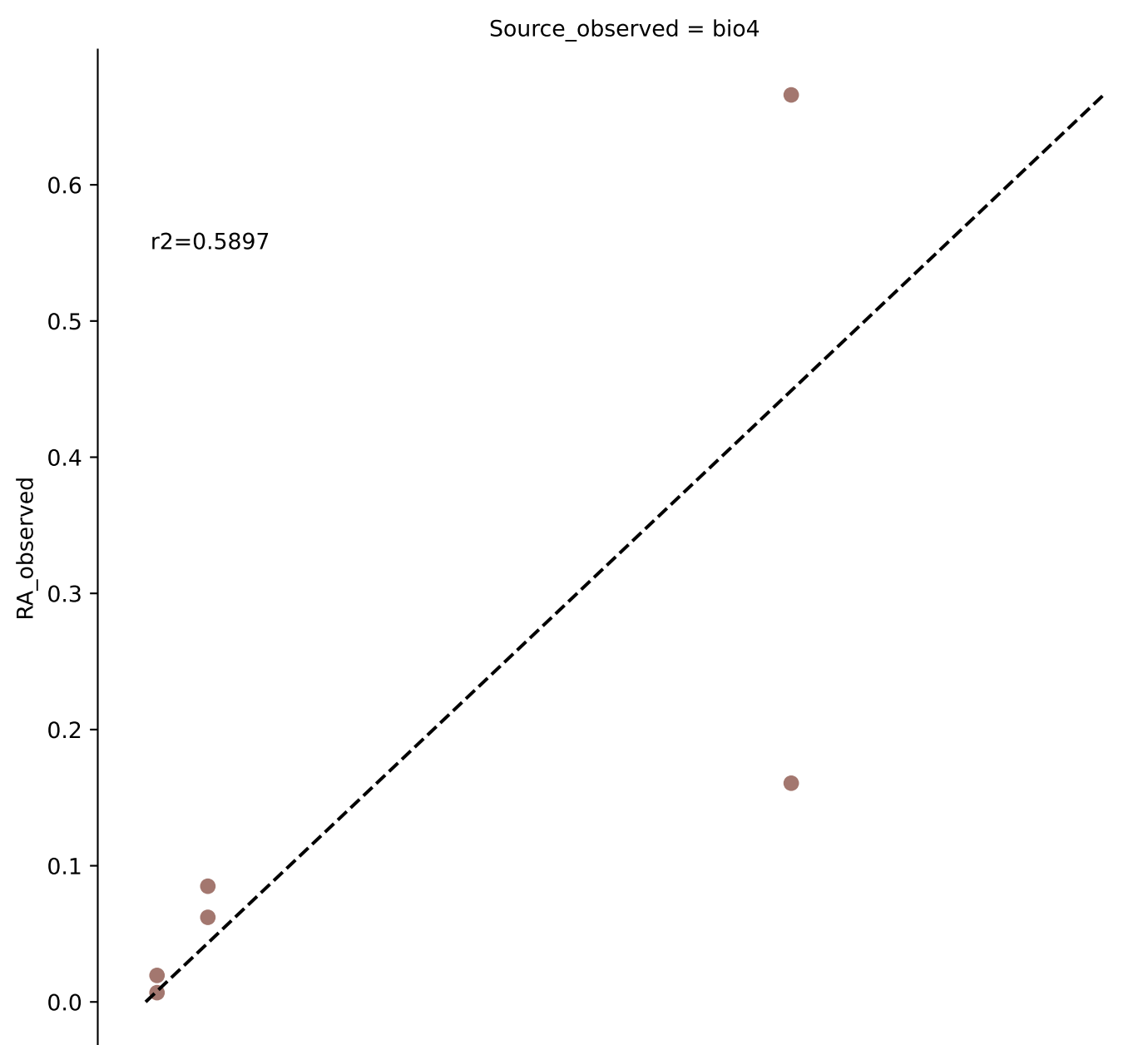


Bivariate Linear Regression for Sample MIX-A in Experiment nist (Species at filter threshold 0)

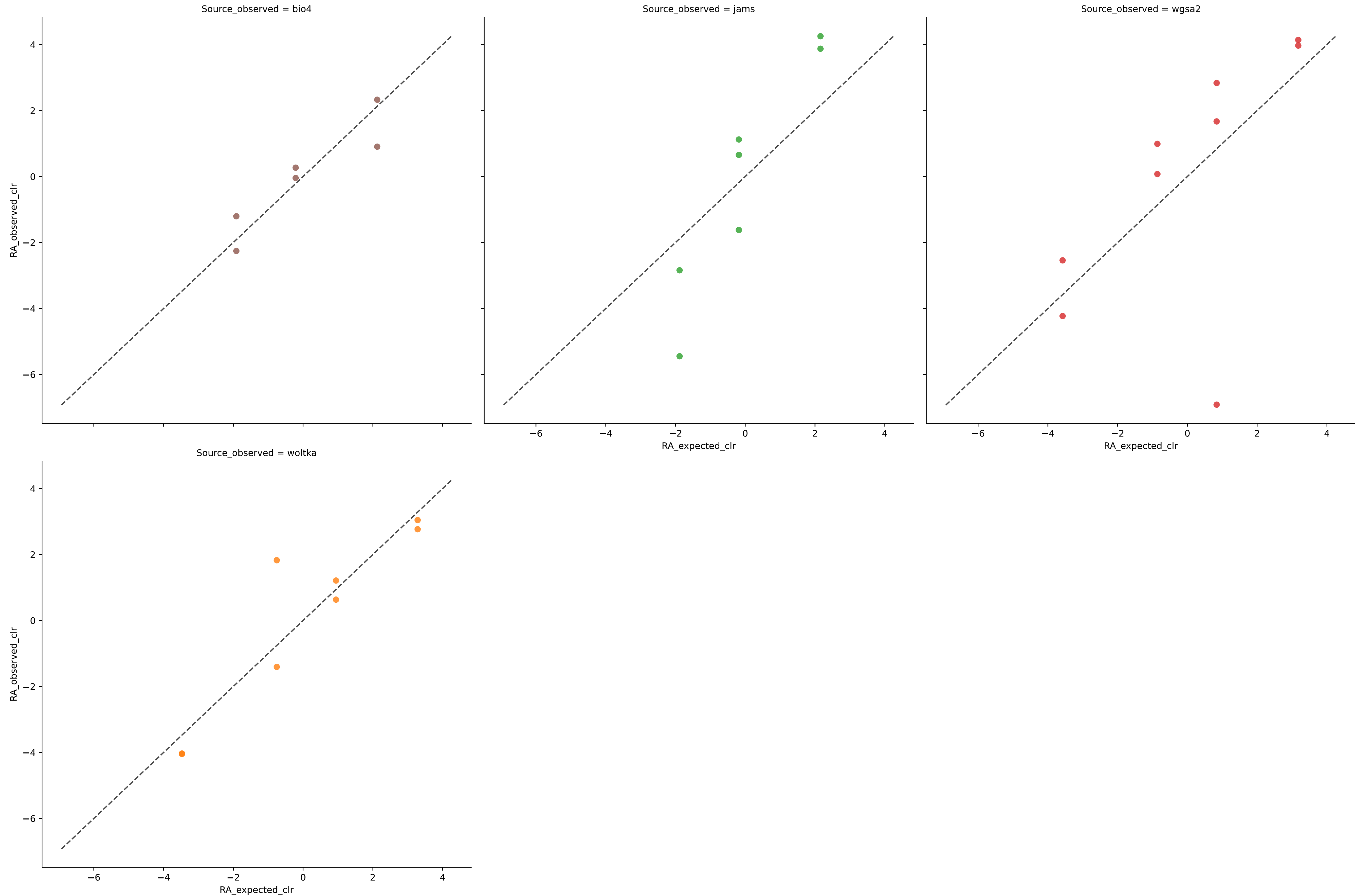


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	7	0.9974	0.0107	1.5752	0.9626	0.0151	100.0000	0.0000
jams	6	0.9765	0.0247	4.3745	0.9258	0.0354	100.0000	0.0000
wgsa2	9	0.5949	0.0720	2.2860	0.6760	0.1369	100.0000	0.0000
woltka	8	0.9172	0.0438	4.1130	0.8249	0.0658	100.0000	0.0000

Bivariate Linear Regression for Sample MIX-B in Experiment nist (Species at filter threshold 0)

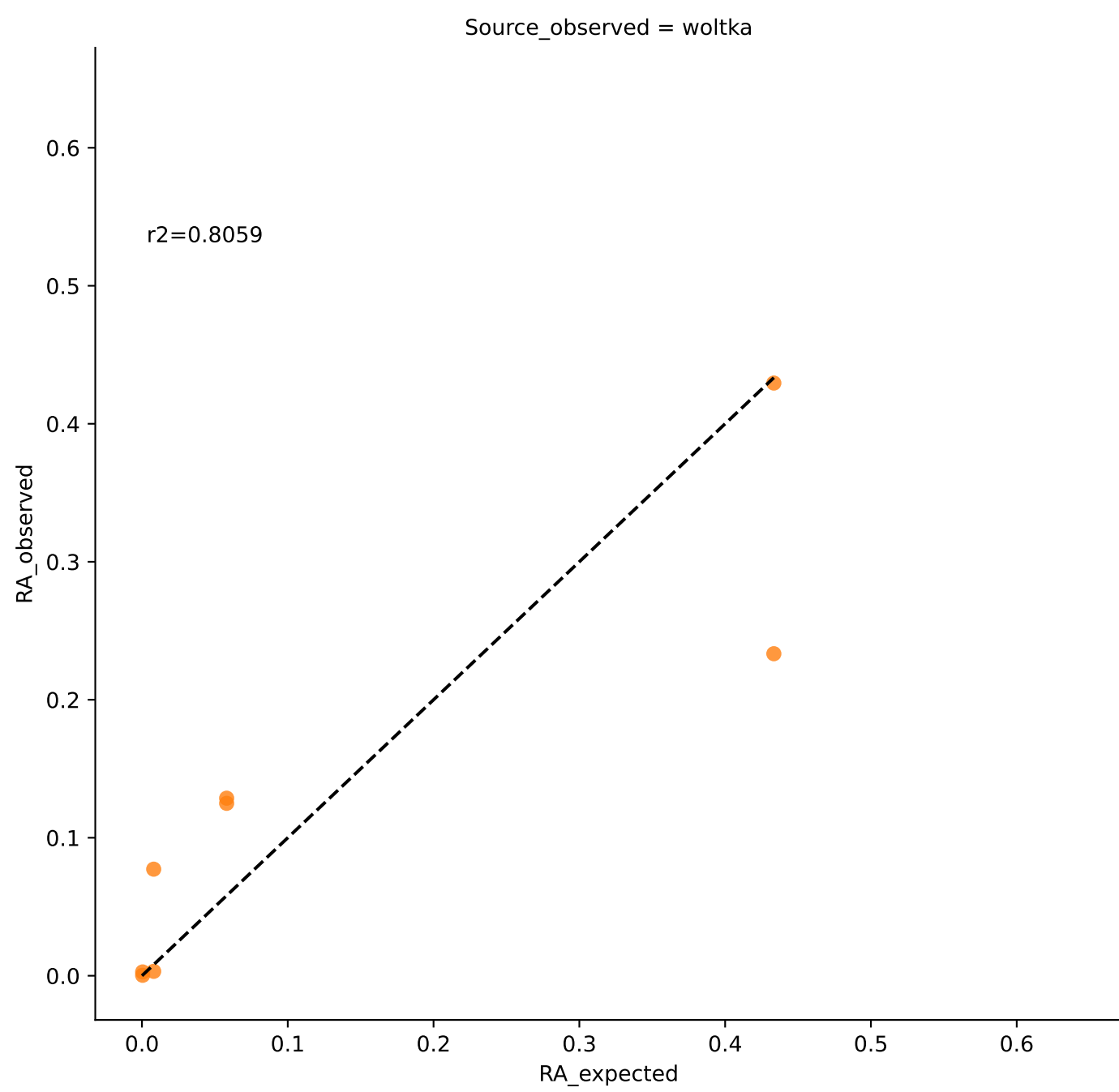
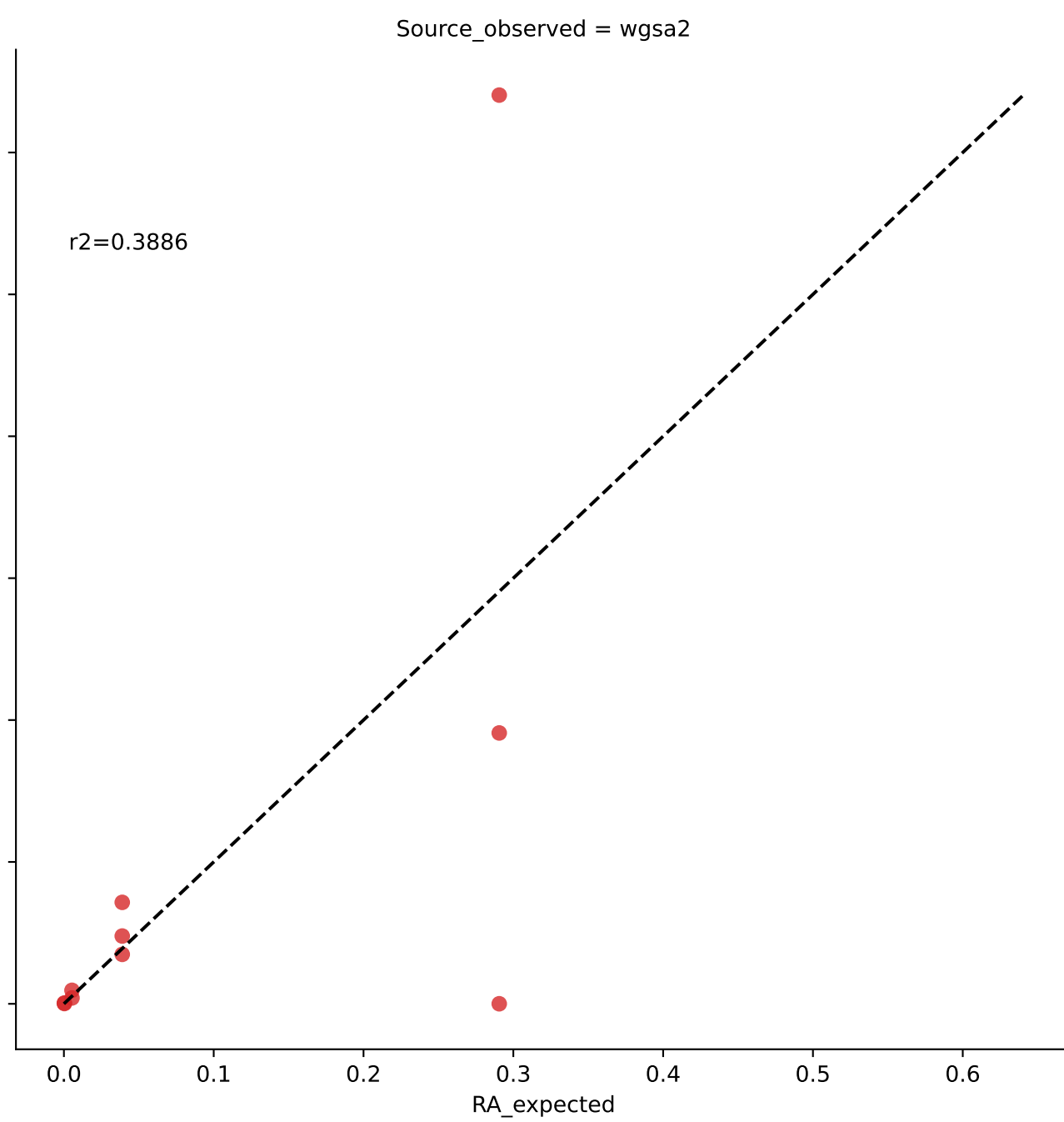
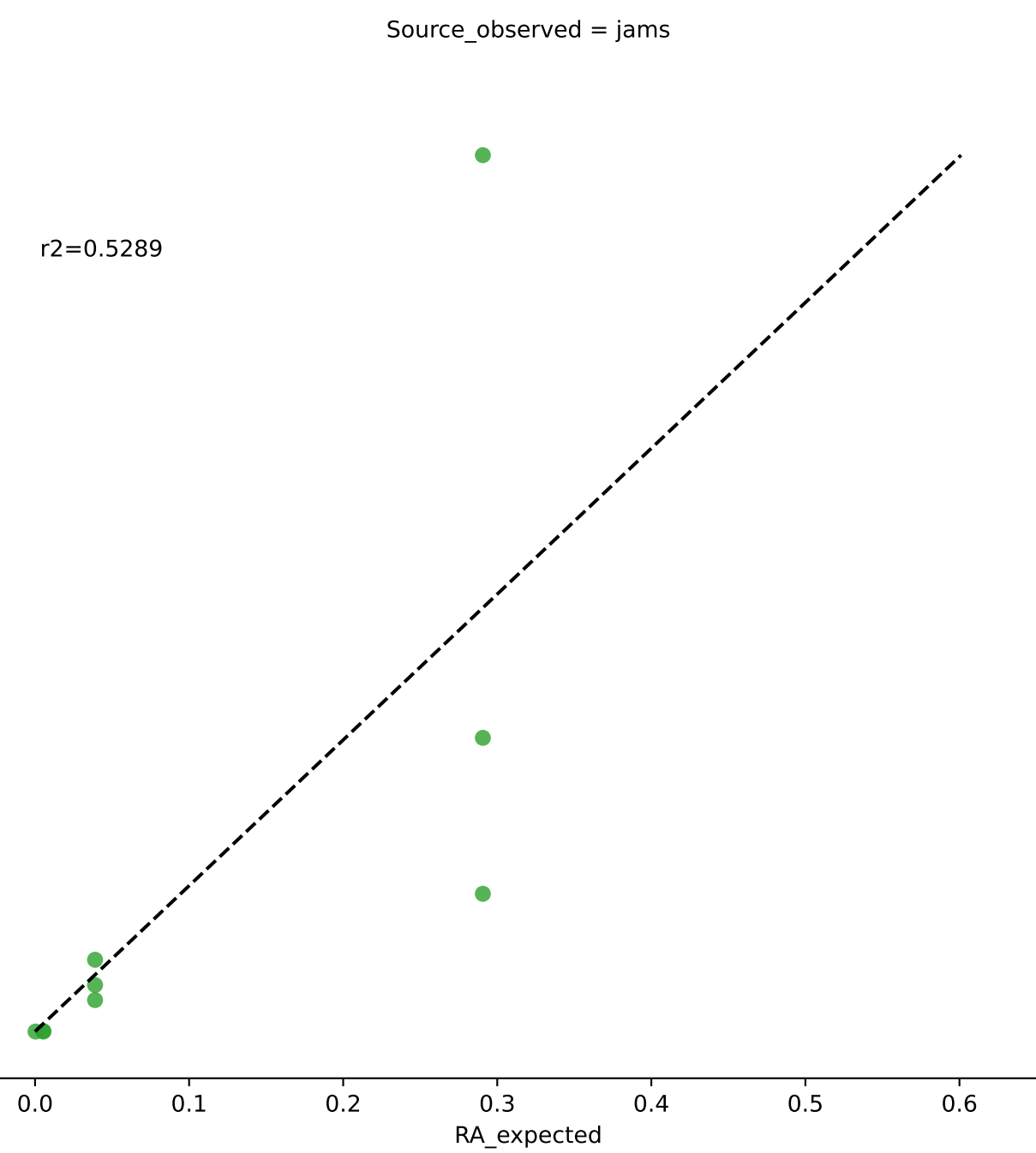
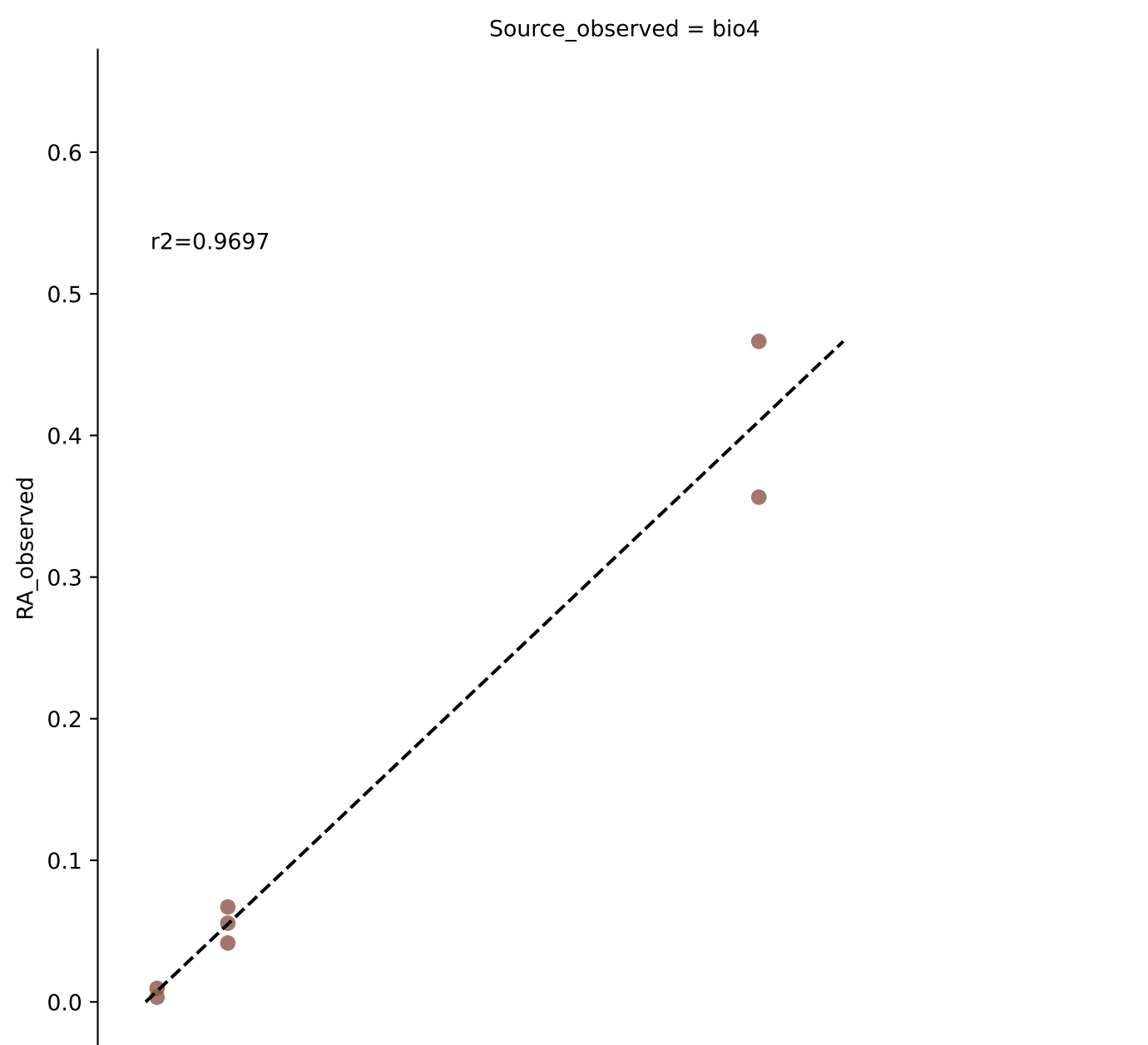


Bivariate Linear Regression for Sample MIX-B in Experiment nist (Species at filter threshold 0)

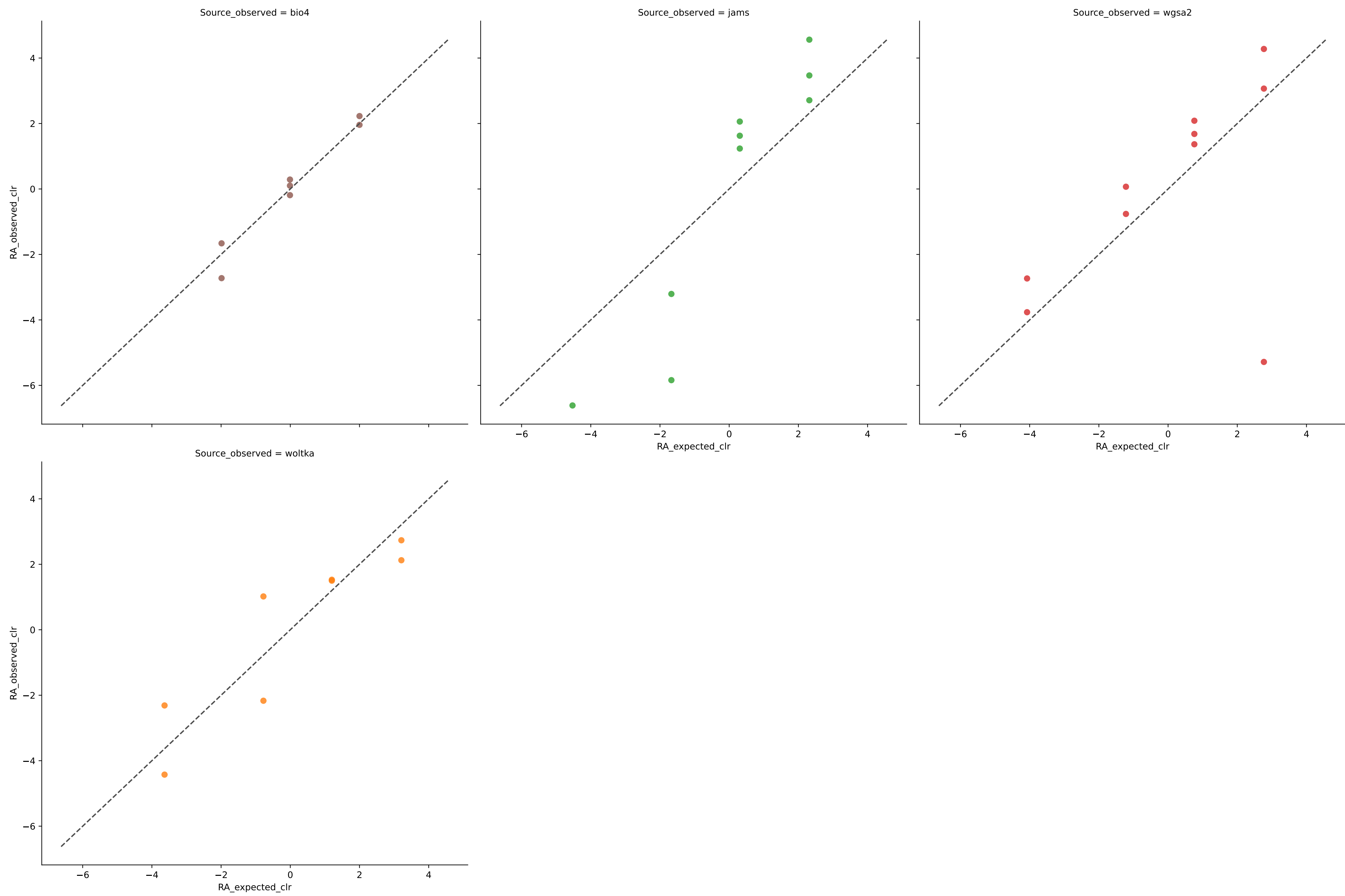


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	6	0.5897	0.0964	1.5530	0.7107	0.1486	100.0000	0.0000
jams	7	0.9484	0.0395	5.0450	0.8619	0.0578	100.0000	0.0000
wgsa2	9	0.9573	0.0228	8.4941	0.8975	0.0356	100.0000	0.0000
woltka	8	0.9096	0.0365	2.8687	0.8542	0.0614	100.0000	0.0000

Bivariate Linear Regression for Sample MIX-C in Experiment nist (Species at filter threshold 0)

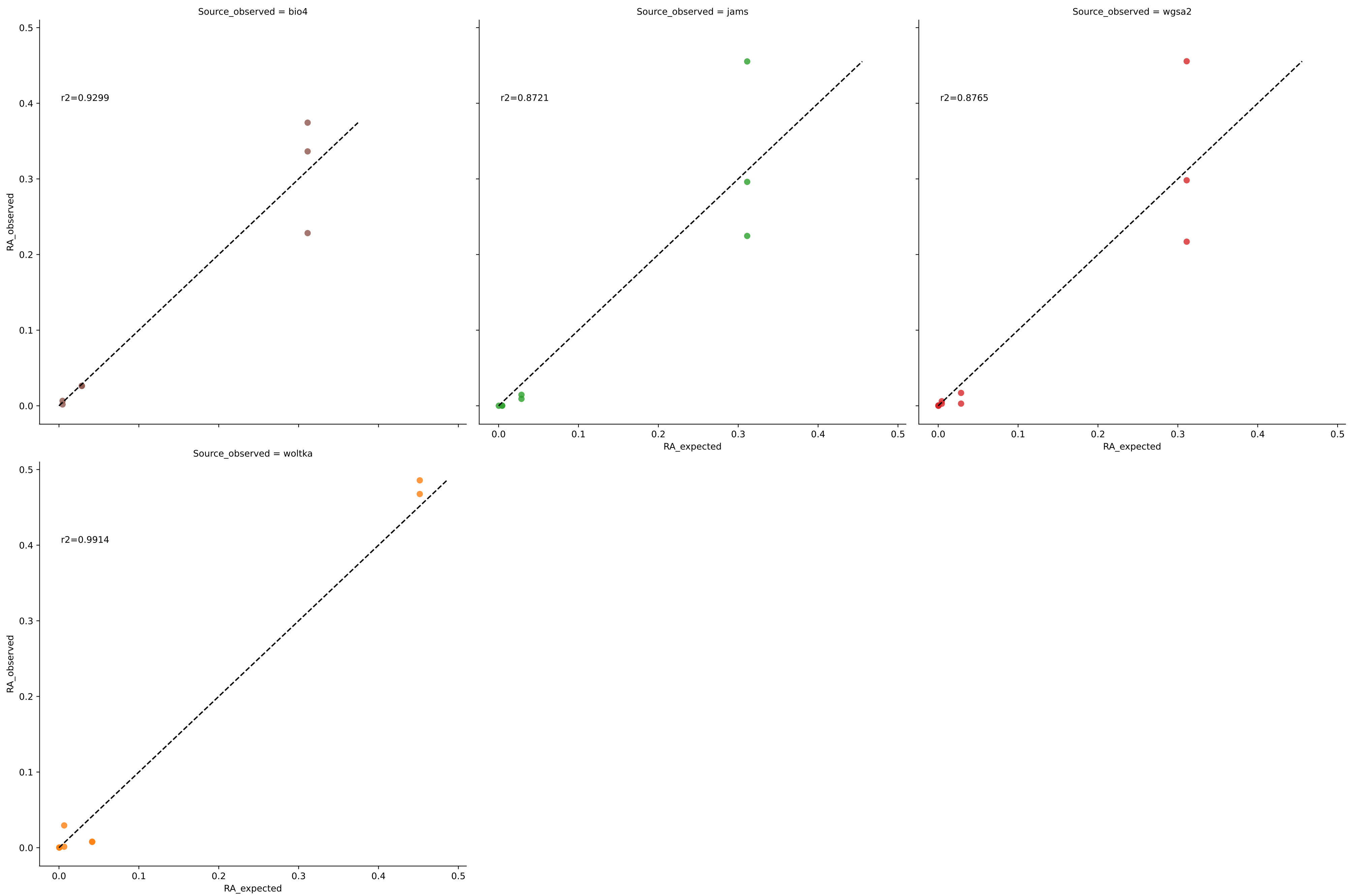


Bivariate Linear Regression for Sample MIX-C in Experiment nist (Species at filter threshold 0)

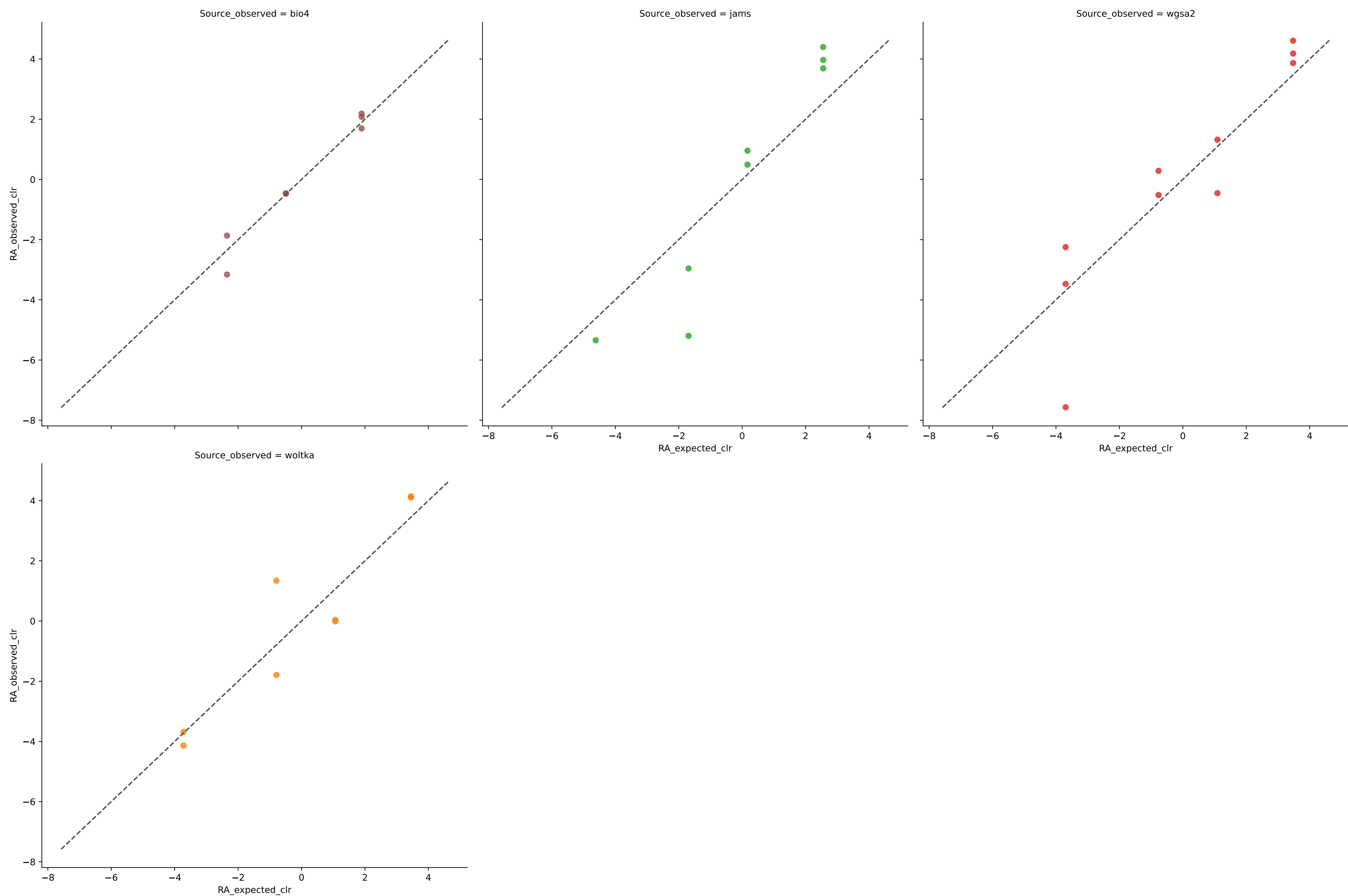


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	7	0.9697	0.0203	0.9148	0.9288	0.0302	100.0000	0.0000
jams	9	0.5289	0.0713	6.0245	0.6793	0.1262	100.0000	0.0000
wgsa2	10	0.3886	0.0791	8.5992	0.6043	0.1476	100.0000	0.0000
woltka	8	0.8059	0.0522	3.0224	0.7911	0.0824	100.0000	0.0000

Bivariate Linear Regression for Sample MIX-D in Experiment nist (Species at filter threshold 0)

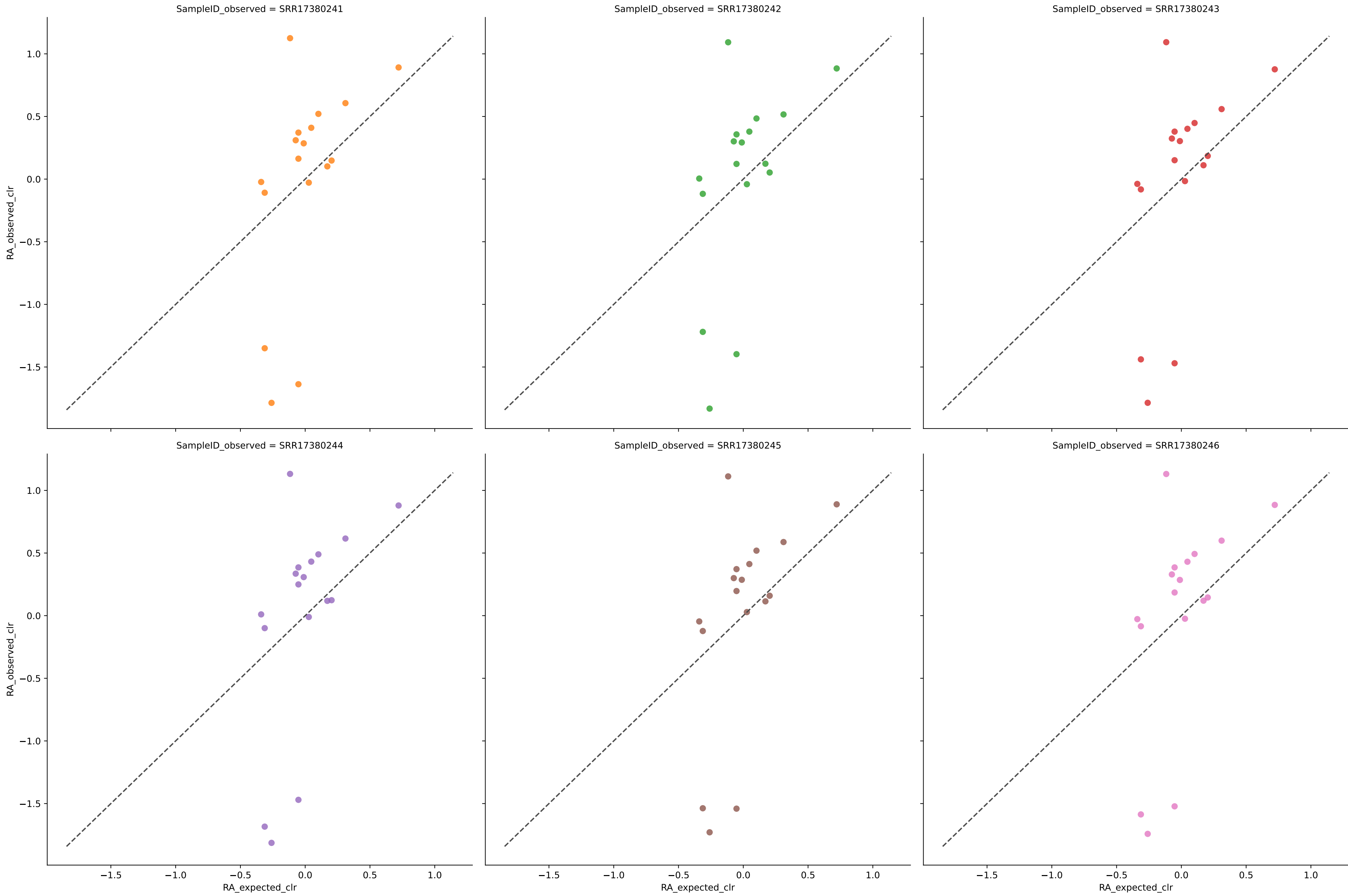


Bivariate Linear Regression for Sample MIX-D in Experiment nist (Species at filter threshold 0)



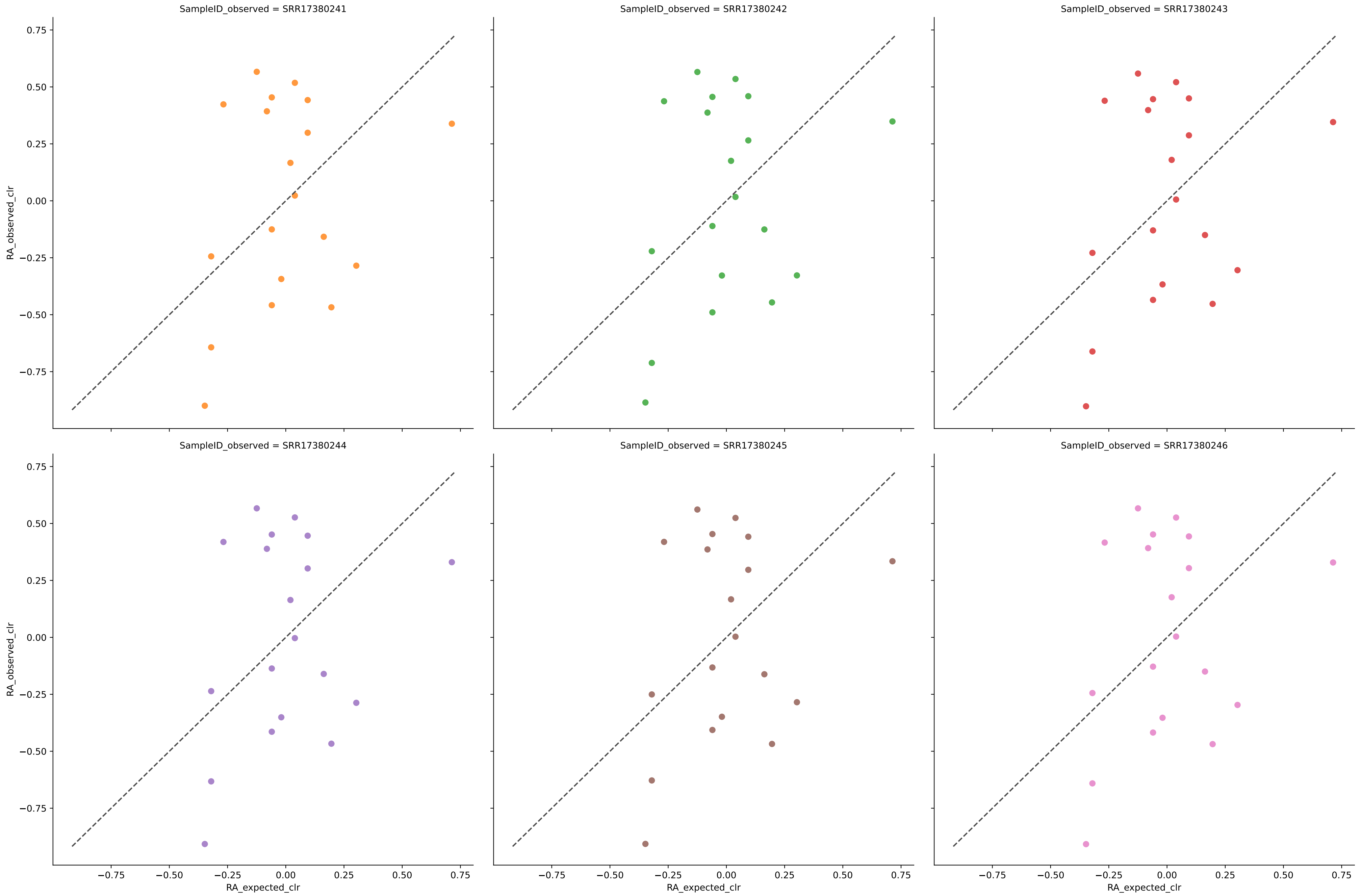
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	7	0.9299	0.0258	1.0255	0.9098	0.0406	100.0000	0.0000
jams	8	0.8721	0.0360	4.6725	0.8559	0.0603	100.0000	0.0000
wgsa2	10	0.8765	0.0293	4.7634	0.8537	0.0554	100.0000	0.0000
woltka	8	0.9914	0.0182	2.9743	0.9270	0.0230	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using bio4 in Experiment tourlousse with filter 0



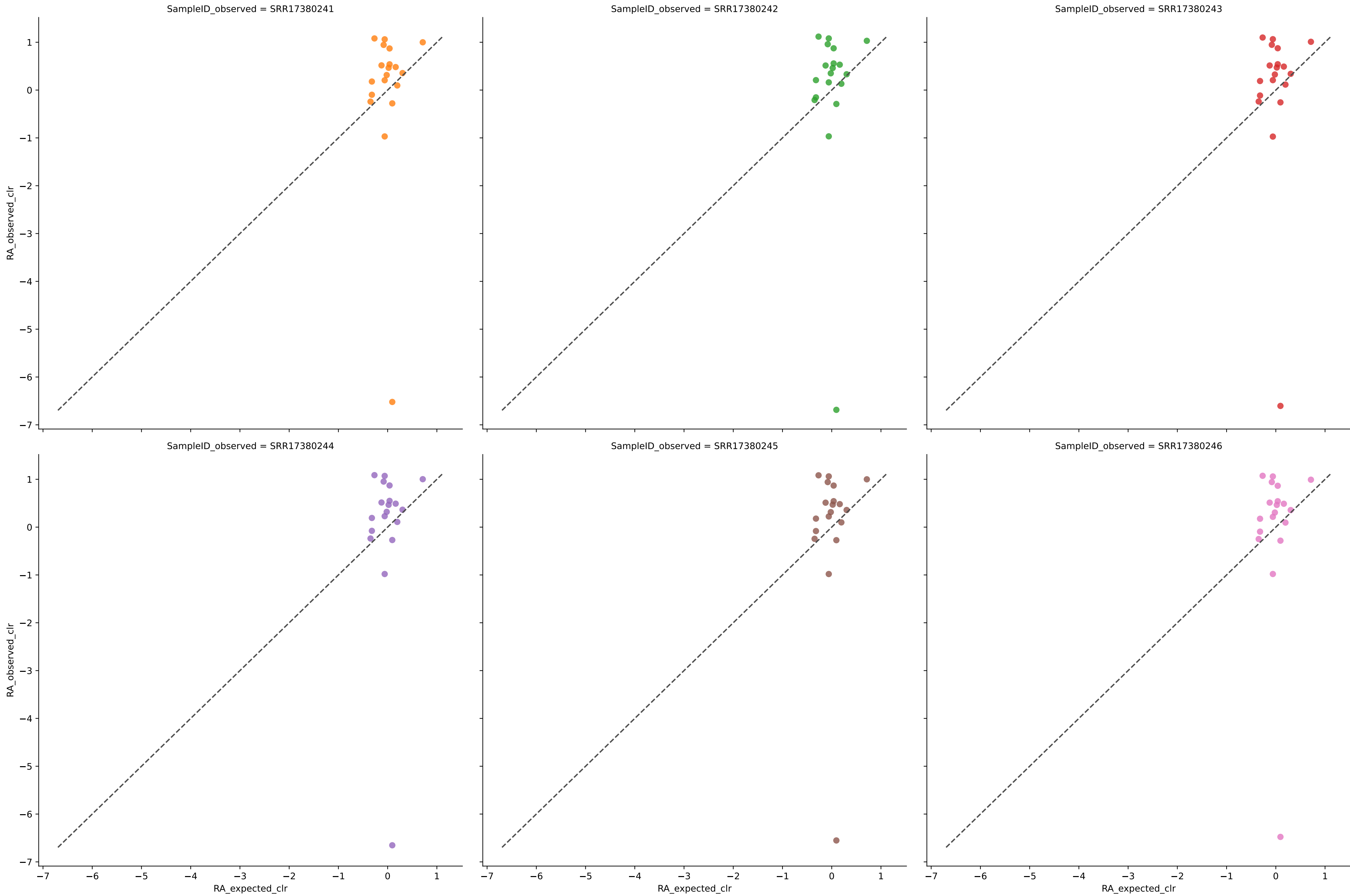
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR17380241	17	0.2683	0.0031	2.9154	0.8404	0.0048	100.0000	0.0000
SRR17380242	17	0.2667	0.0030	2.7401	0.8446	0.0047	100.0000	0.0000
SRR17380243	17	0.2744	0.0030	2.8350	0.8461	0.0046	100.0000	0.0000
SRR17380244	17	0.2551	0.0032	3.0019	0.8349	0.0048	100.0000	0.0000
SRR17380245	17	0.2771	0.0031	2.8973	0.8429	0.0047	100.0000	0.0000
SRR17380246	17	0.2608	0.0032	2.9331	0.8386	0.0048	100.0000	0.0000
Average	17	0.2671	0.0031	2.8871	0.8412	0.0048	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using jams in Experiment tourlousse with filter 0



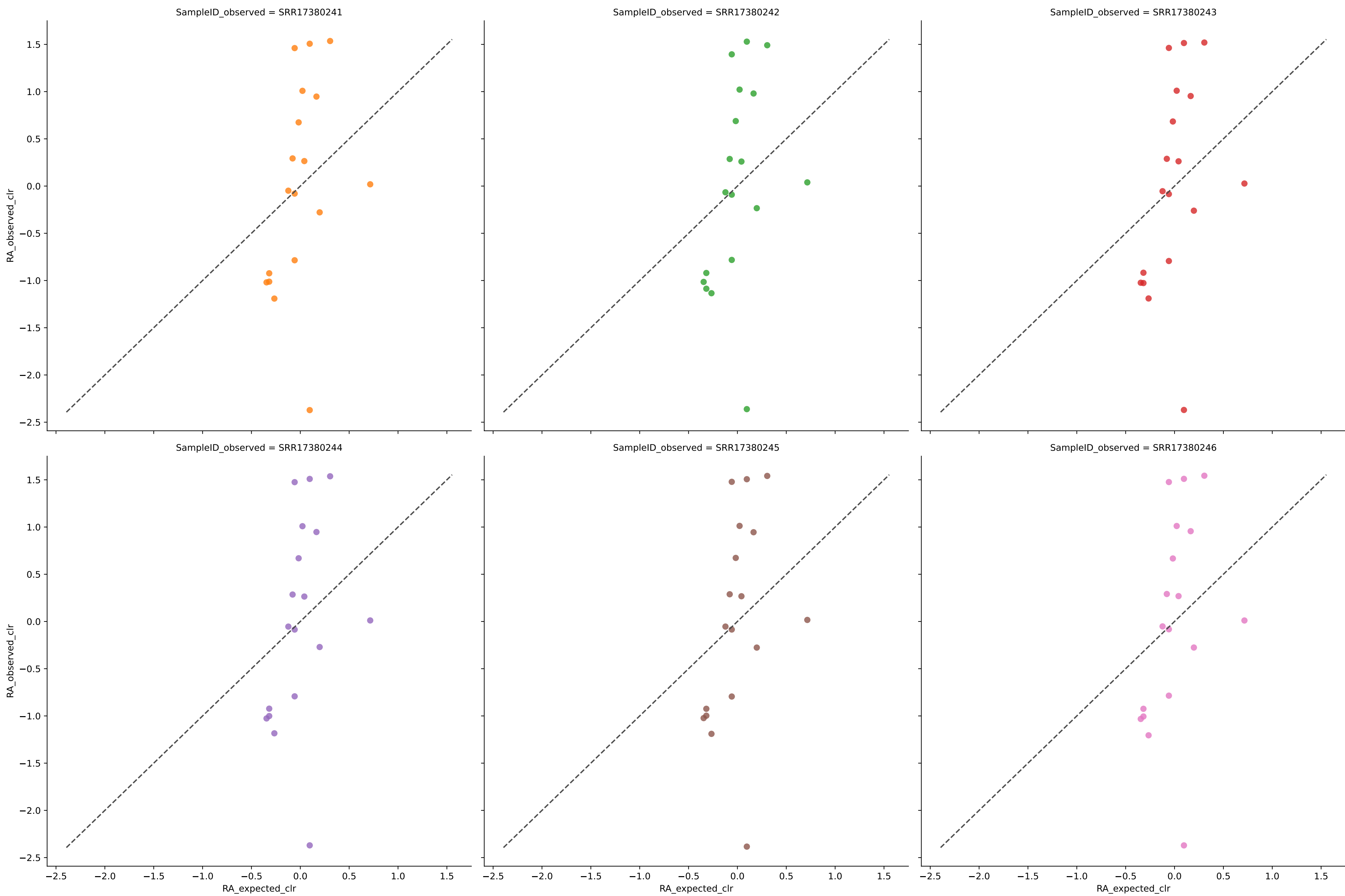
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR17380241	19	0.0294	0.0033	1.8893	0.8104	0.0039	100.0000	0.0000
SRR17380242	19	0.0298	0.0033	1.9110	0.8093	0.0039	100.0000	0.0000
SRR17380243	19	0.0296	0.0033	1.8964	0.8091	0.0039	100.0000	0.0000
SRR17380244	19	0.0276	0.0033	1.8859	0.8093	0.0039	100.0000	0.0000
SRR17380245	19	0.0291	0.0033	1.8776	0.8108	0.0039	100.0000	0.0000
SRR17380246	19	0.0276	0.0033	1.8902	0.8094	0.0039	100.0000	0.0000
Average	19	0.0289	0.0033	1.8917	0.8097	0.0039	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using wgsa in Experiment tourlousse with filter 0



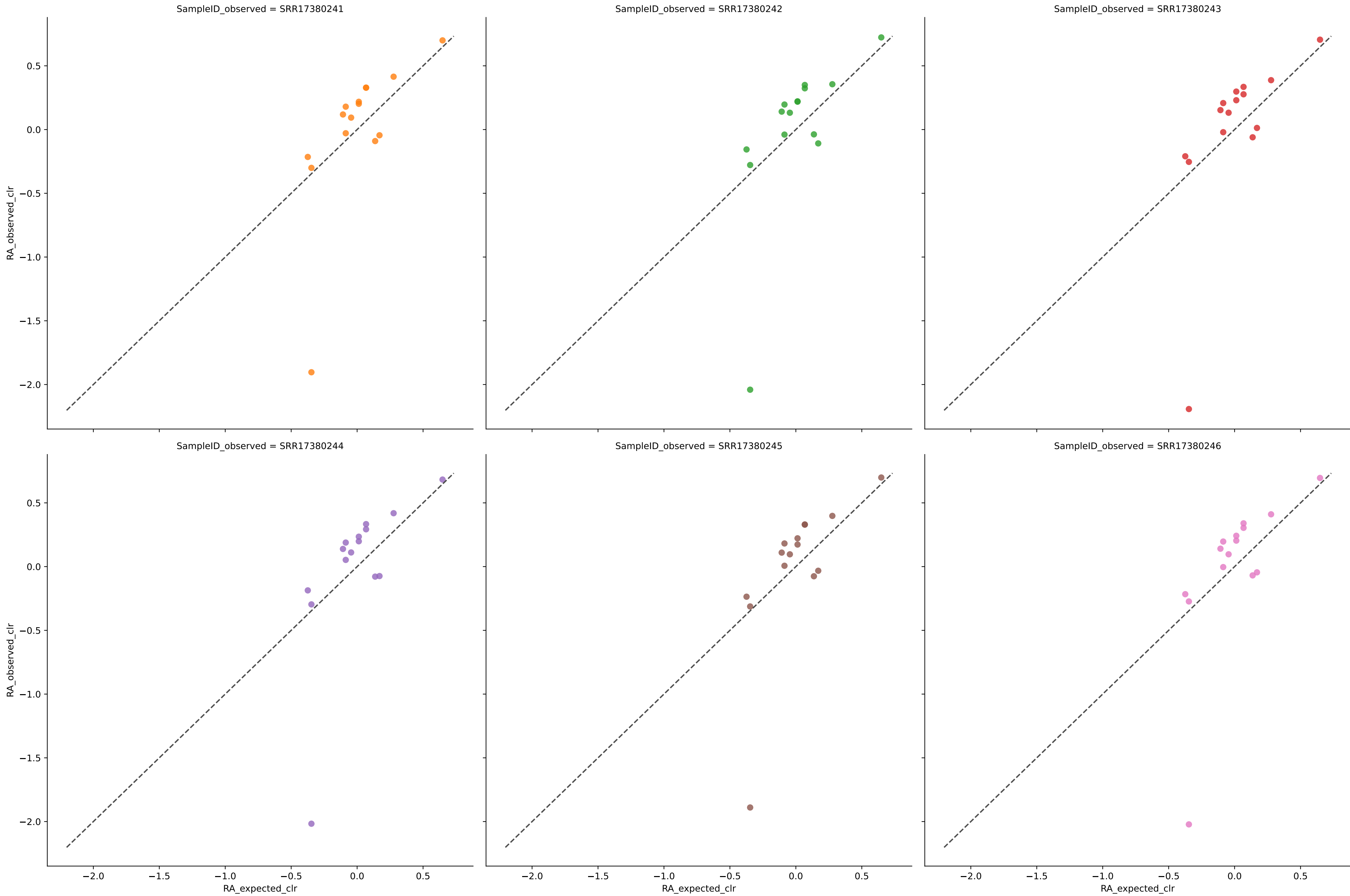
	Diversity	R ²	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR17380241	19	0.0476	0.0036	7.1479	0.7940	0.0048	100.0000	0.0000
SRR17380242	19	0.0495	0.0036	7.3192	0.7927	0.0048	100.0000	0.0000
SRR17380243	19	0.0477	0.0036	7.2308	0.7938	0.0048	100.0000	0.0000
SRR17380244	19	0.0464	0.0036	7.2784	0.7946	0.0048	100.0000	0.0000
SRR17380245	19	0.0474	0.0036	7.1787	0.7948	0.0048	100.0000	0.0000
SRR17380246	19	0.0471	0.0036	7.1059	0.7944	0.0048	100.0000	0.0000
Average	19	0.0476	0.0036	7.2102	0.7941	0.0048	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using woltka in Experiment tourlousse with filter 0



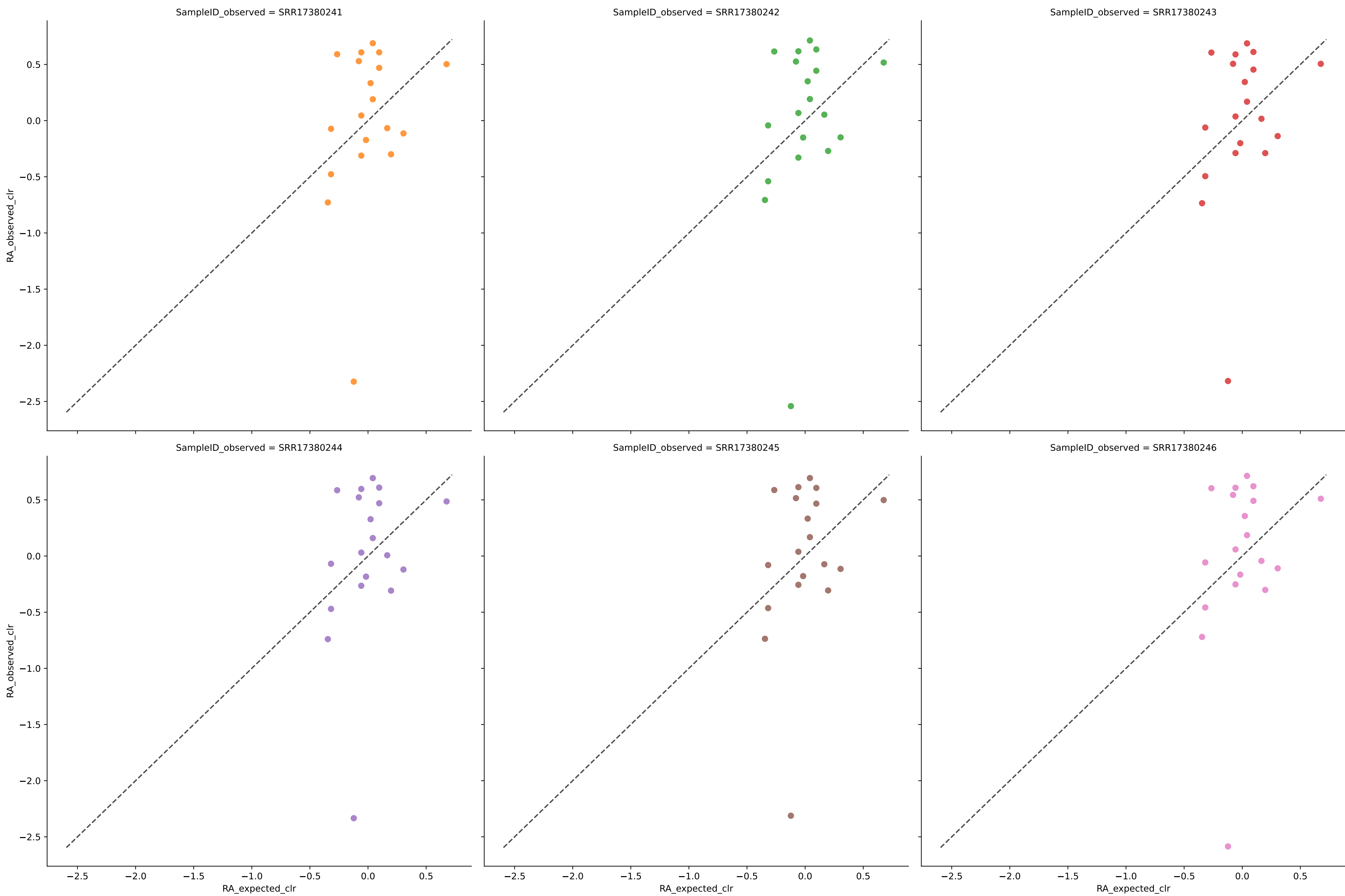
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR17380241	18	0.0619	0.0067	4.1905	0.6387	0.0082	100.0000	0.0000
SRR17380242	18	0.0655	0.0066	4.1572	0.6414	0.0081	100.0000	0.0000
SRR17380243	18	0.0618	0.0067	4.1904	0.6385	0.0082	100.0000	0.0000
SRR17380244	18	0.0608	0.0067	4.1939	0.6374	0.0083	100.0000	0.0000
SRR17380245	18	0.0609	0.0067	4.2048	0.6373	0.0083	100.0000	0.0000
SRR17380246	18	0.0611	0.0067	4.2050	0.6370	0.0083	100.0000	0.0000
Average	18	0.0620	0.0067	4.1903	0.6384	0.0082	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using bio4 in Experiment tourlousse with filter 0



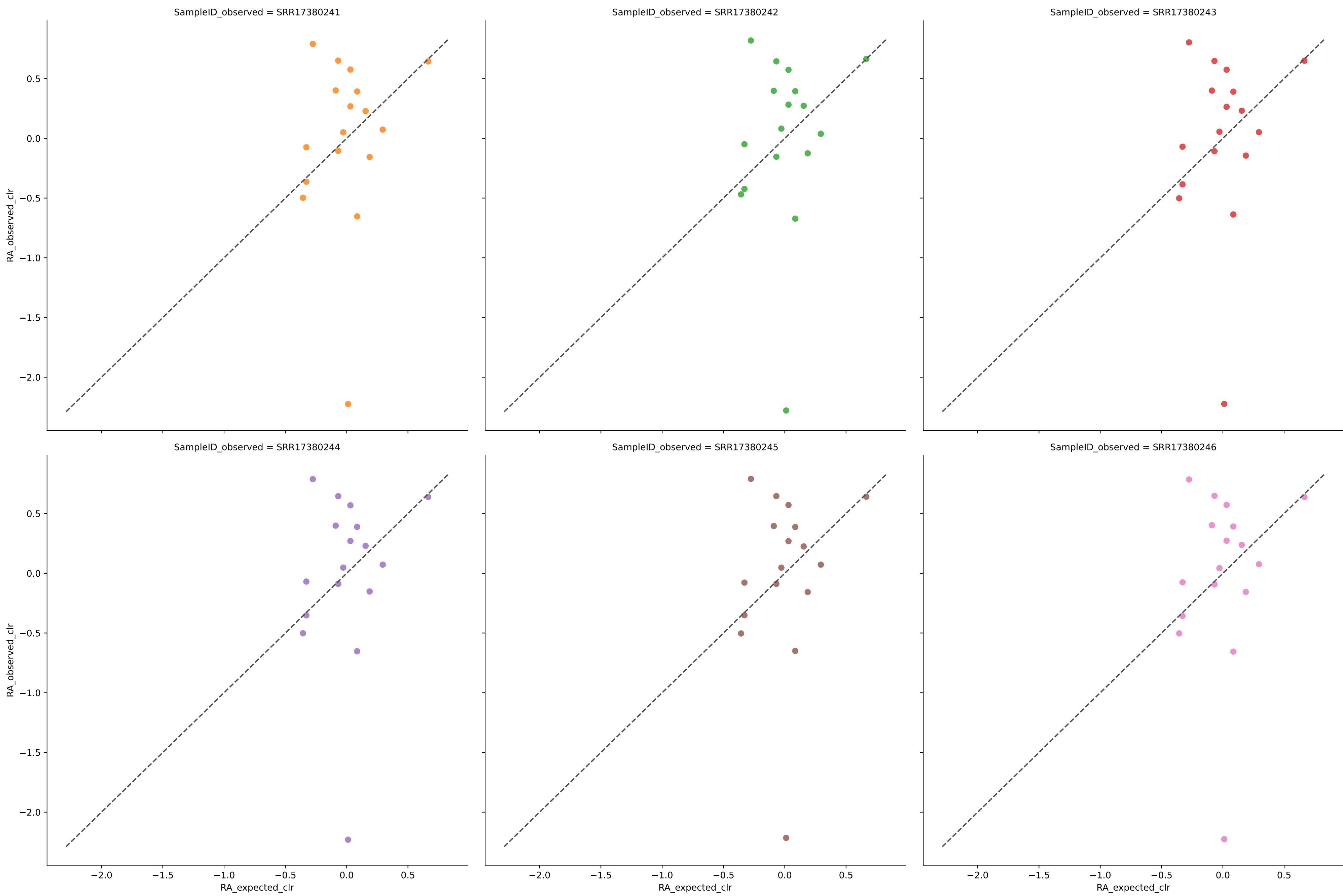
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR17380241	15	0.6588	0.0018	1.7116	0.9196	0.0023	100.0000	0.0000
SRR17380242	15	0.6349	0.0018	1.8567	0.9185	0.0024	100.0000	0.0000
SRR17380243	15	0.6416	0.0018	1.9887	0.9203	0.0024	100.0000	0.0000
SRR17380244	15	0.6284	0.0019	1.8273	0.9153	0.0024	100.0000	0.0000
SRR17380245	15	0.6673	0.0017	1.6908	0.9221	0.0023	100.0000	0.0000
SRR17380246	15	0.6450	0.0018	1.8257	0.9199	0.0024	100.0000	0.0000
Average	15	0.6460	0.0018	1.8168	0.9193	0.0024	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using jams in Experiment tourlousse with filter 0



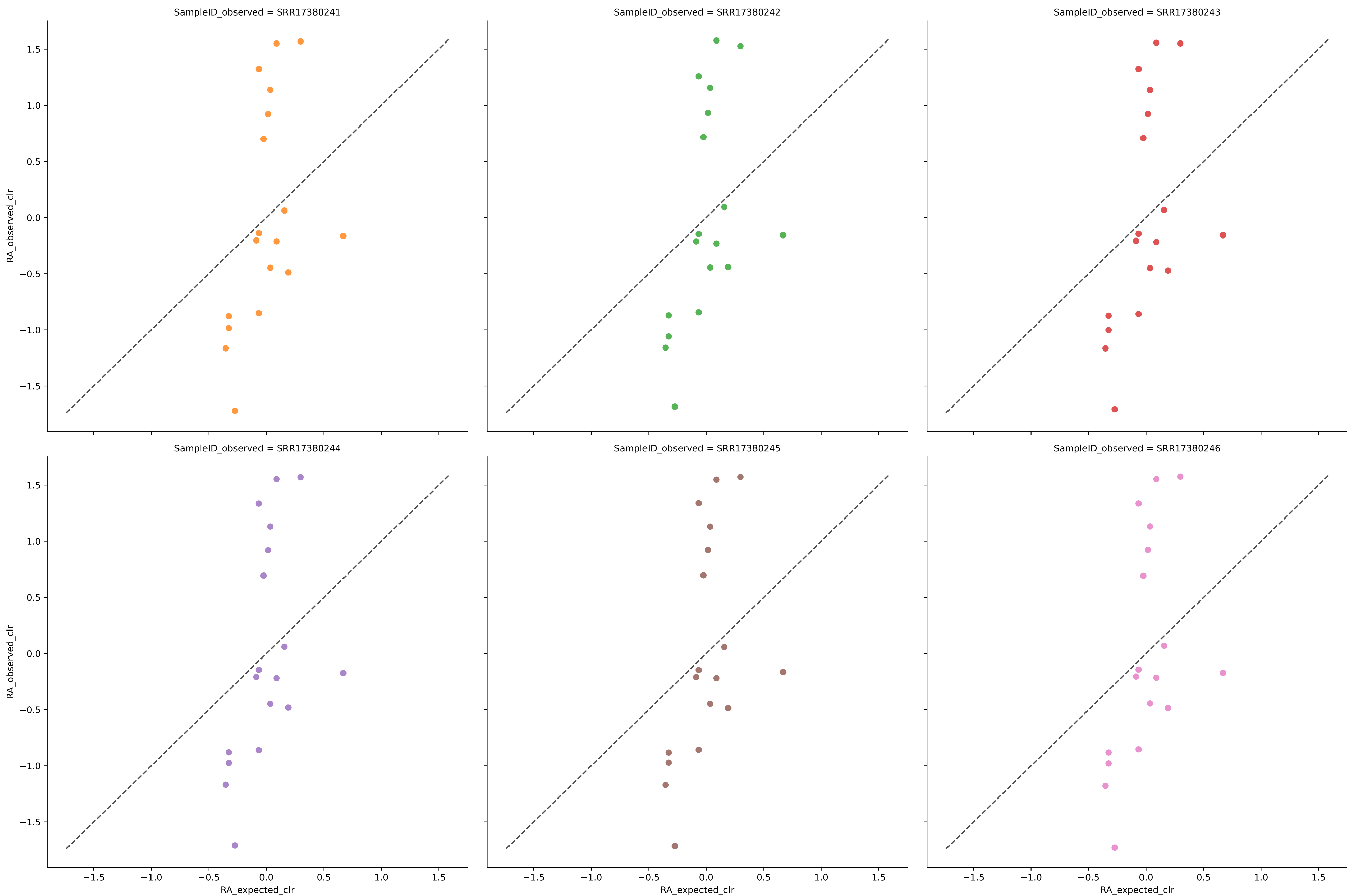
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR17380241	19	0.0685	0.0035	2.8585	0.8028	0.0040	100.0000	0.0000
SRR17380242	19	0.0699	0.0034	3.0443	0.8034	0.0041	100.0000	0.0000
SRR17380243	19	0.0715	0.0034	2.8464	0.8049	0.0040	100.0000	0.0000
SRR17380244	19	0.0671	0.0034	2.8565	0.8044	0.0040	100.0000	0.0000
SRR17380245	19	0.0674	0.0034	2.8435	0.8038	0.0040	100.0000	0.0000
SRR17380246	19	0.0661	0.0035	3.0762	0.8022	0.0041	100.0000	0.0000
Average	19	0.0684	0.0034	2.9209	0.8036	0.0040	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using wgsa in Experiment tourlousse with filter 0



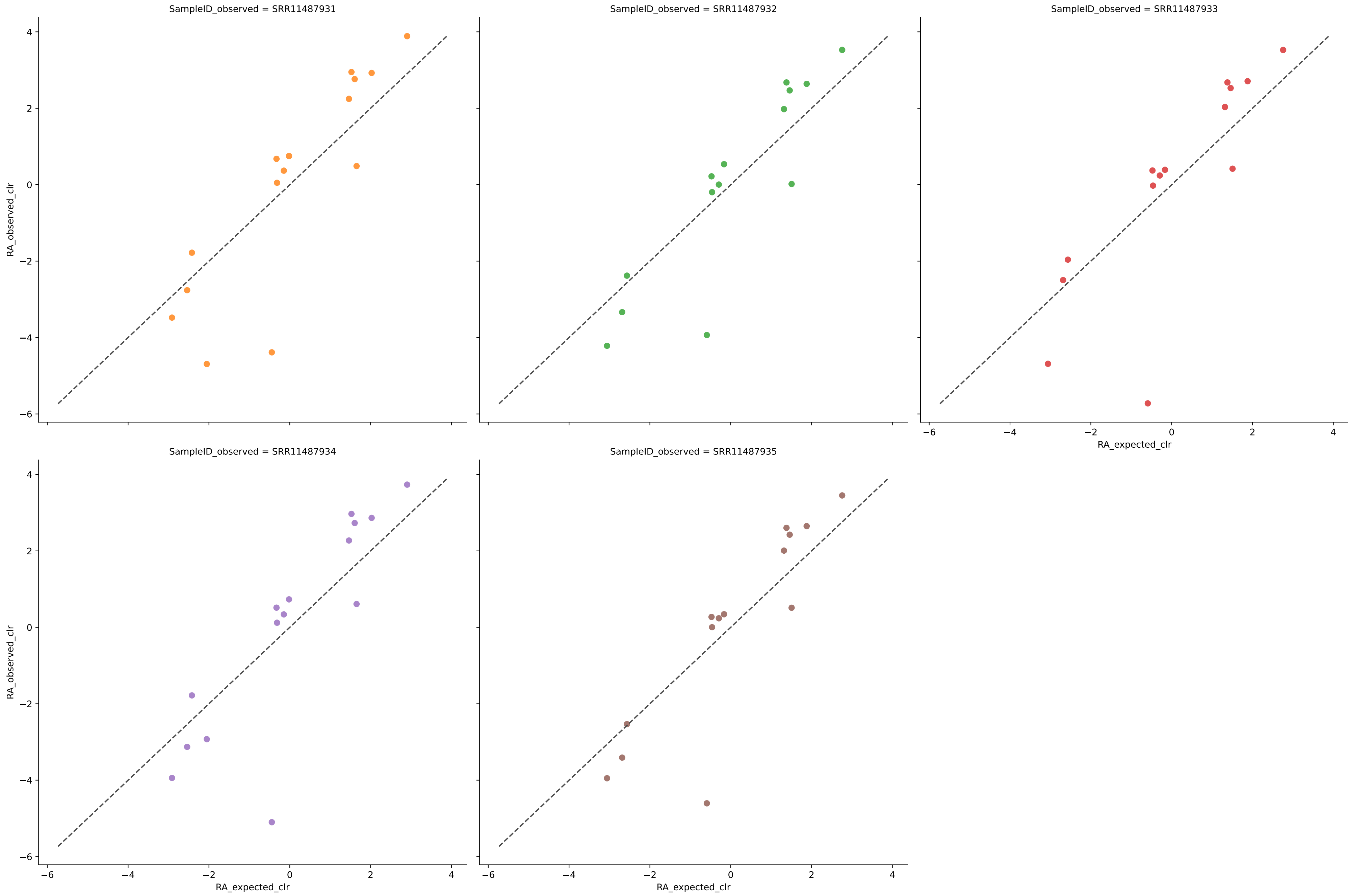
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR17380241	17	0.0588	0.0036	2.8562	0.8157	0.0047	100.0000	0.0000
SRR17380242	17	0.0603	0.0036	2.9180	0.8144	0.0047	100.0000	0.0000
SRR17380243	17	0.0586	0.0036	2.8552	0.8154	0.0047	100.0000	0.0000
SRR17380244	17	0.0579	0.0036	2.8562	0.8166	0.0047	100.0000	0.0000
SRR17380245	17	0.0583	0.0036	2.8440	0.8167	0.0047	100.0000	0.0000
SRR17380246	17	0.0586	0.0036	2.8549	0.8162	0.0047	100.0000	0.0000
Average	17	0.0587	0.0036	2.8641	0.8158	0.0047	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using woltka in Experiment tourlousse with filter 0



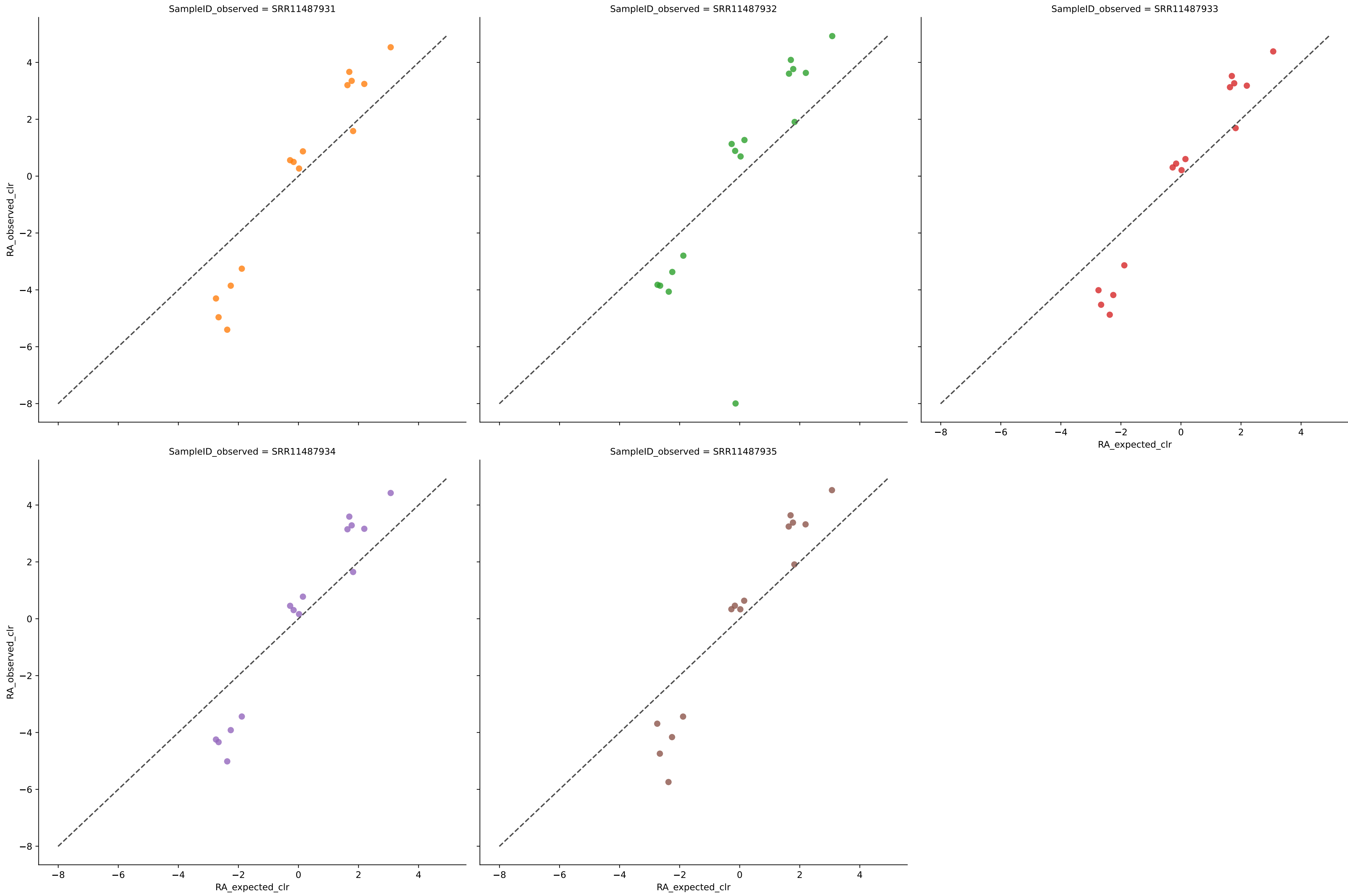
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR17380241	18	0.0560	0.0073	3.7226	0.6054	0.0086	100.0000	0.0000
SRR17380242	18	0.0568	0.0072	3.6942	0.6082	0.0084	100.0000	0.0000
SRR17380243	18	0.0555	0.0073	3.7162	0.6058	0.0086	100.0000	0.0000
SRR17380244	18	0.0550	0.0073	3.7255	0.6044	0.0086	100.0000	0.0000
SRR17380245	18	0.0554	0.0073	3.7281	0.6042	0.0086	100.0000	0.0000
SRR17380246	18	0.0555	0.0073	3.7368	0.6044	0.0086	100.0000	0.0000
Average	18	0.0557	0.0073	3.7206	0.6054	0.0086	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using bio4 in Experiment Amos hilo with filter 0



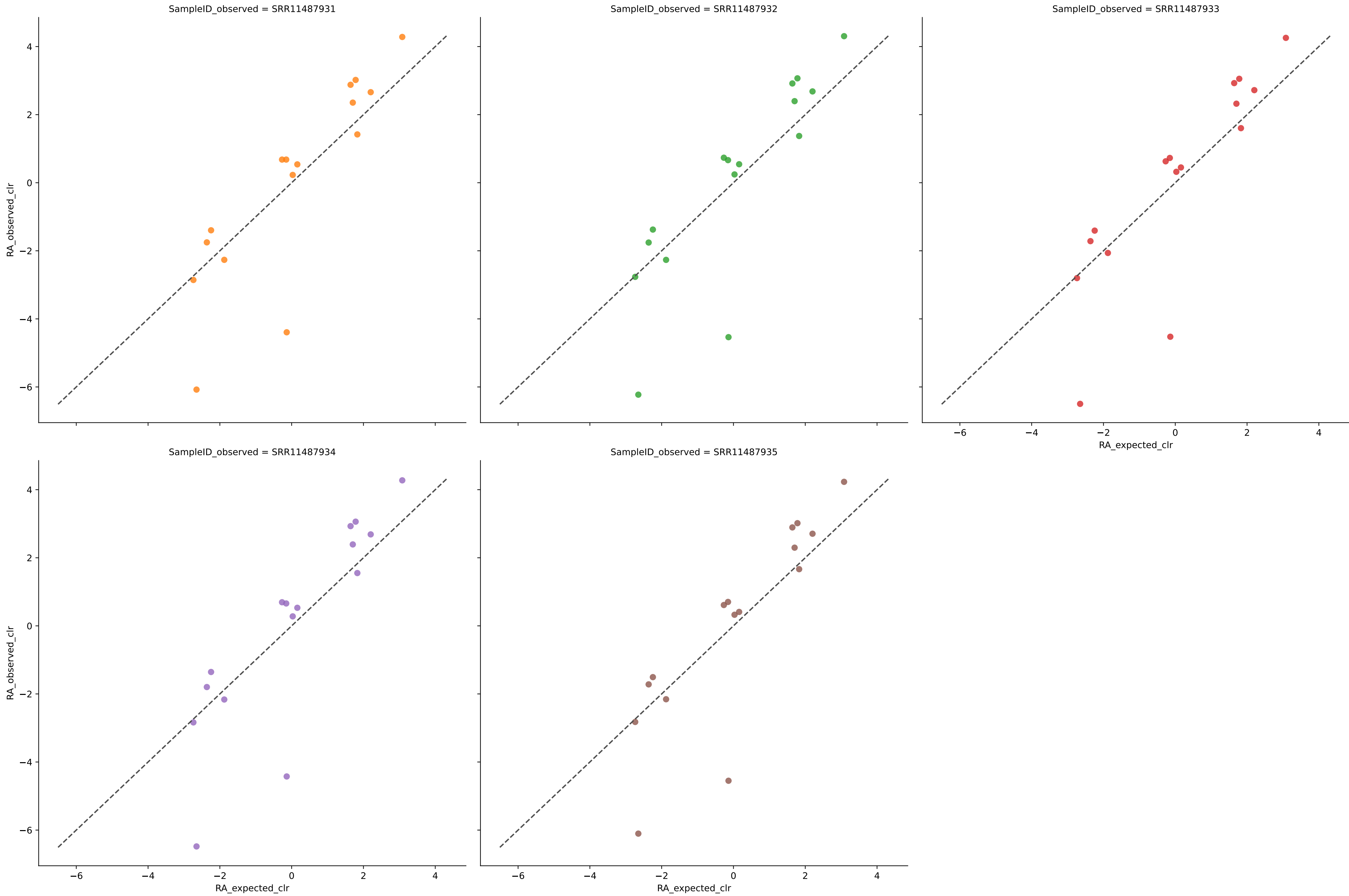
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487931	15	0.9163	0.0035	5.6916	0.8671	0.0064	100.0000	0.0000
SRR11487932	14	0.9020	0.0037	4.5410	0.8713	0.0067	100.0000	0.0000
SRR11487933	14	0.9056	0.0034	6.0565	0.8801	0.0064	100.0000	0.0000
SRR11487934	15	0.8969	0.0032	5.6939	0.8806	0.0065	100.0000	0.0000
SRR11487935	14	0.9105	0.0032	4.8690	0.8881	0.0061	100.0000	0.0000
Average	14	0.9063	0.0034	5.3704	0.8774	0.0064	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using jams in Experiment Amos hilo with filter 0



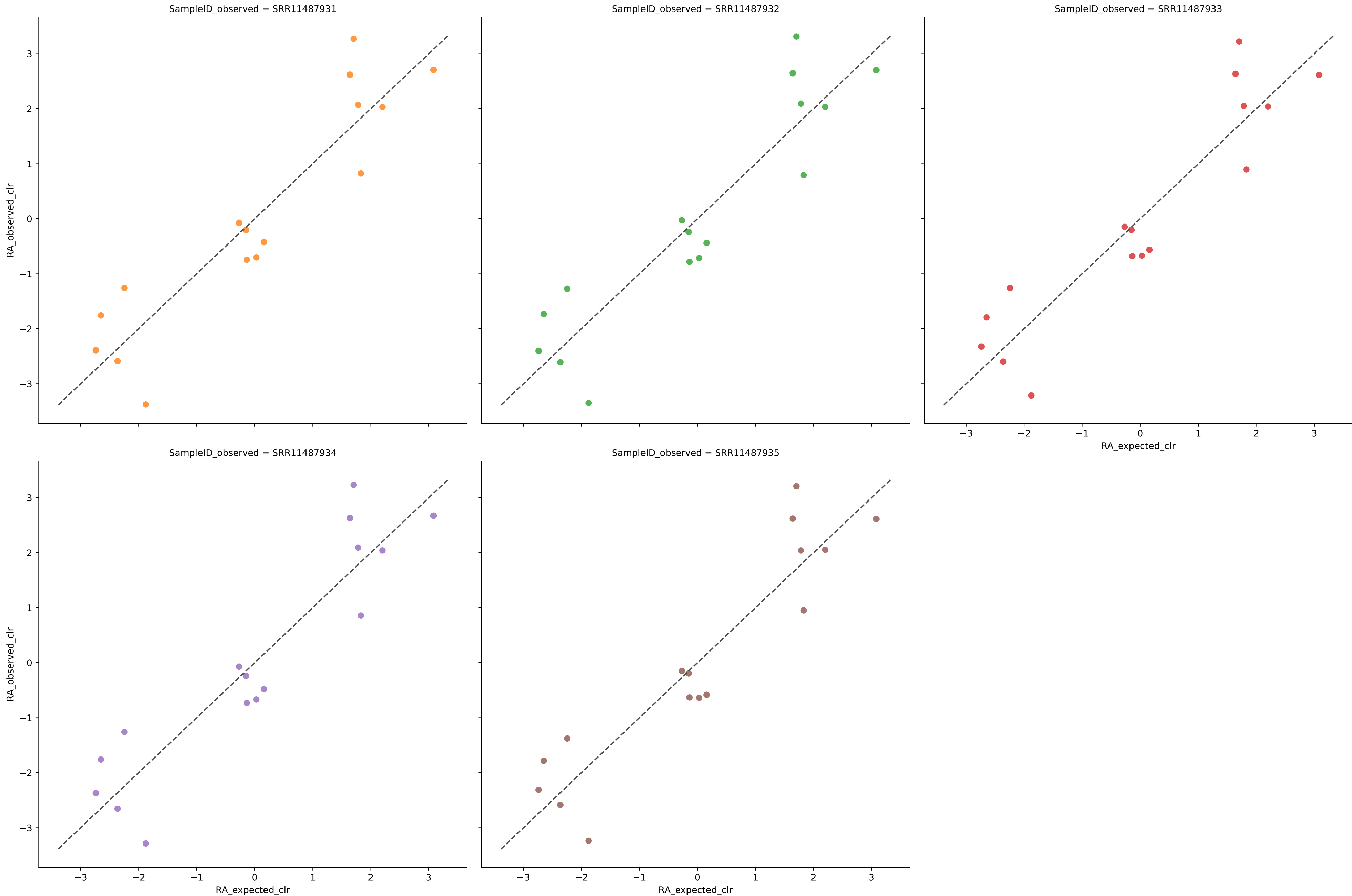
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487931	15	0.9019	0.0045	5.9316	0.8303	0.0071	100.0000	0.0000
SRR11487932	16	0.8973	0.0045	9.6443	0.8223	0.0070	100.0000	0.0000
SRR11487933	15	0.9096	0.0043	5.2991	0.8402	0.0066	100.0000	0.0000
SRR11487934	15	0.9006	0.0044	5.4209	0.8332	0.0069	100.0000	0.0000
SRR11487935	15	0.9155	0.0042	6.0090	0.8419	0.0064	100.0000	0.0000
Average	15	0.9050	0.0044	6.4610	0.8336	0.0068	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using wgsa in Experiment Amos hilo with filter 0



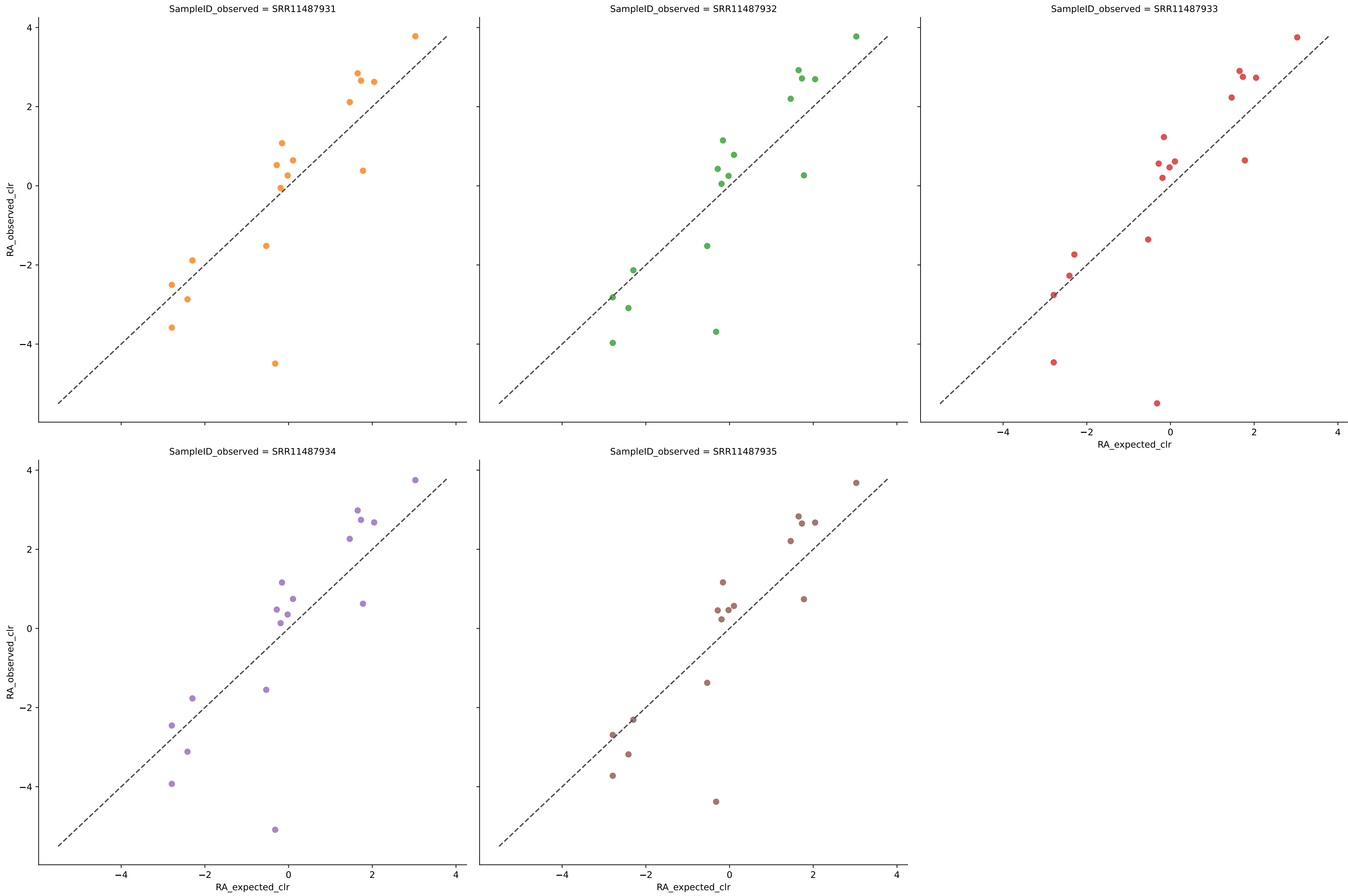
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487931	16	0.9134	0.0048	6.1797	0.8063	0.0083	100.0000	0.0000
SRR11487932	16	0.9112	0.0049	6.4008	0.8046	0.0082	100.0000	0.0000
SRR11487933	16	0.9198	0.0046	6.5033	0.8142	0.0076	100.0000	0.0000
SRR11487934	16	0.9165	0.0047	6.4448	0.8117	0.0078	100.0000	0.0000
SRR11487935	16	0.9235	0.0046	6.2562	0.8179	0.0075	100.0000	0.0000
Average	16	0.9169	0.0047	6.3570	0.8109	0.0079	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using wol in Experiment Amos hilo with filter 0



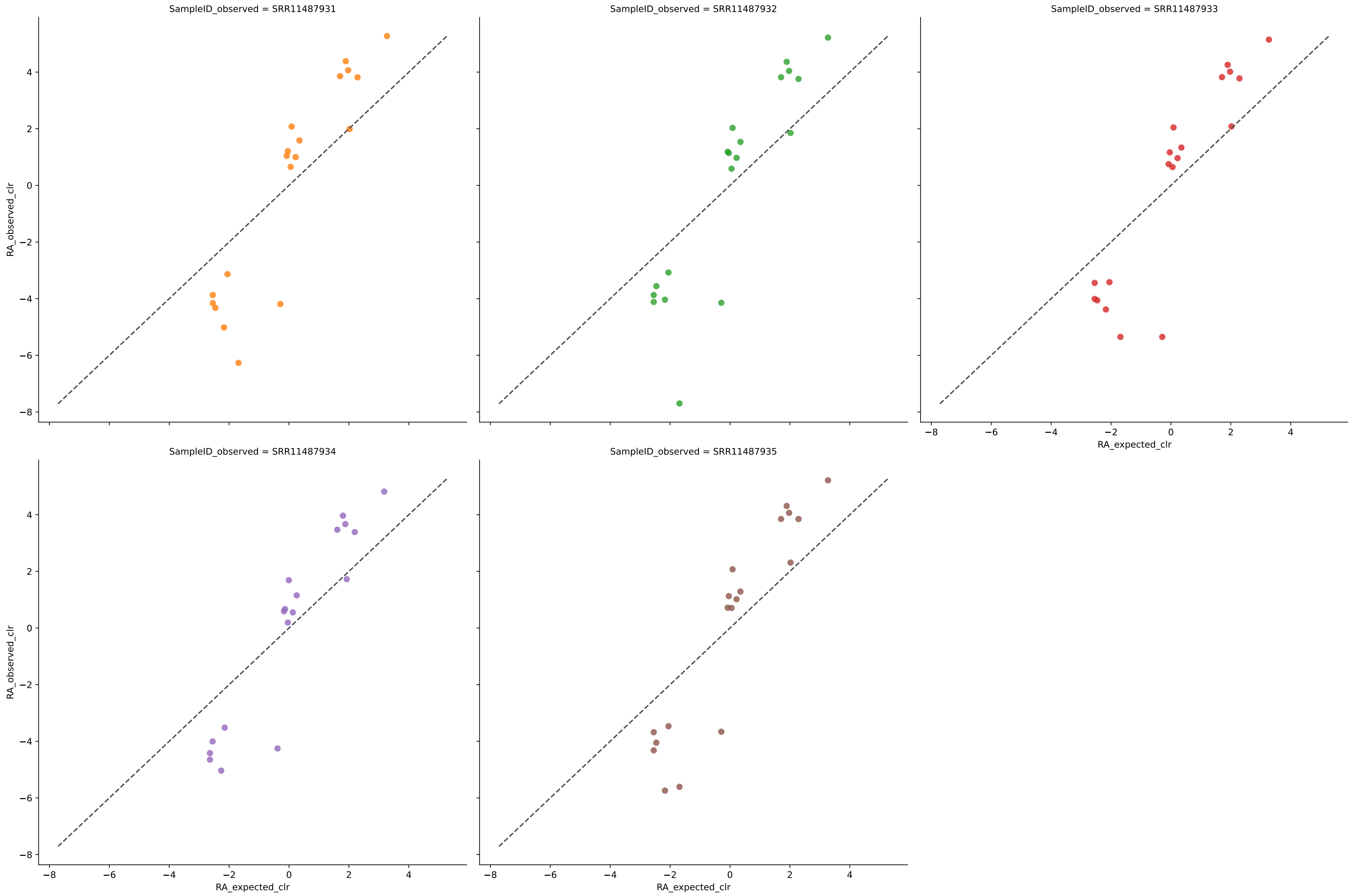
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487931	16	0.3723	0.0086	3.1882	0.6543	0.0166	100.0000	0.0000
SRR11487932	16	0.3532	0.0088	3.2368	0.6454	0.0171	100.0000	0.0000
SRR11487933	16	0.3586	0.0087	3.0849	0.6509	0.0168	100.0000	0.0000
SRR11487934	16	0.3760	0.0086	3.1331	0.6548	0.0165	100.0000	0.0000
SRR11487935	16	0.3660	0.0086	3.0234	0.6564	0.0166	100.0000	0.0000
Average	16	0.3652	0.0087	3.1333	0.6524	0.0167	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using bio4 in Experiment Amos hilo with filter 0



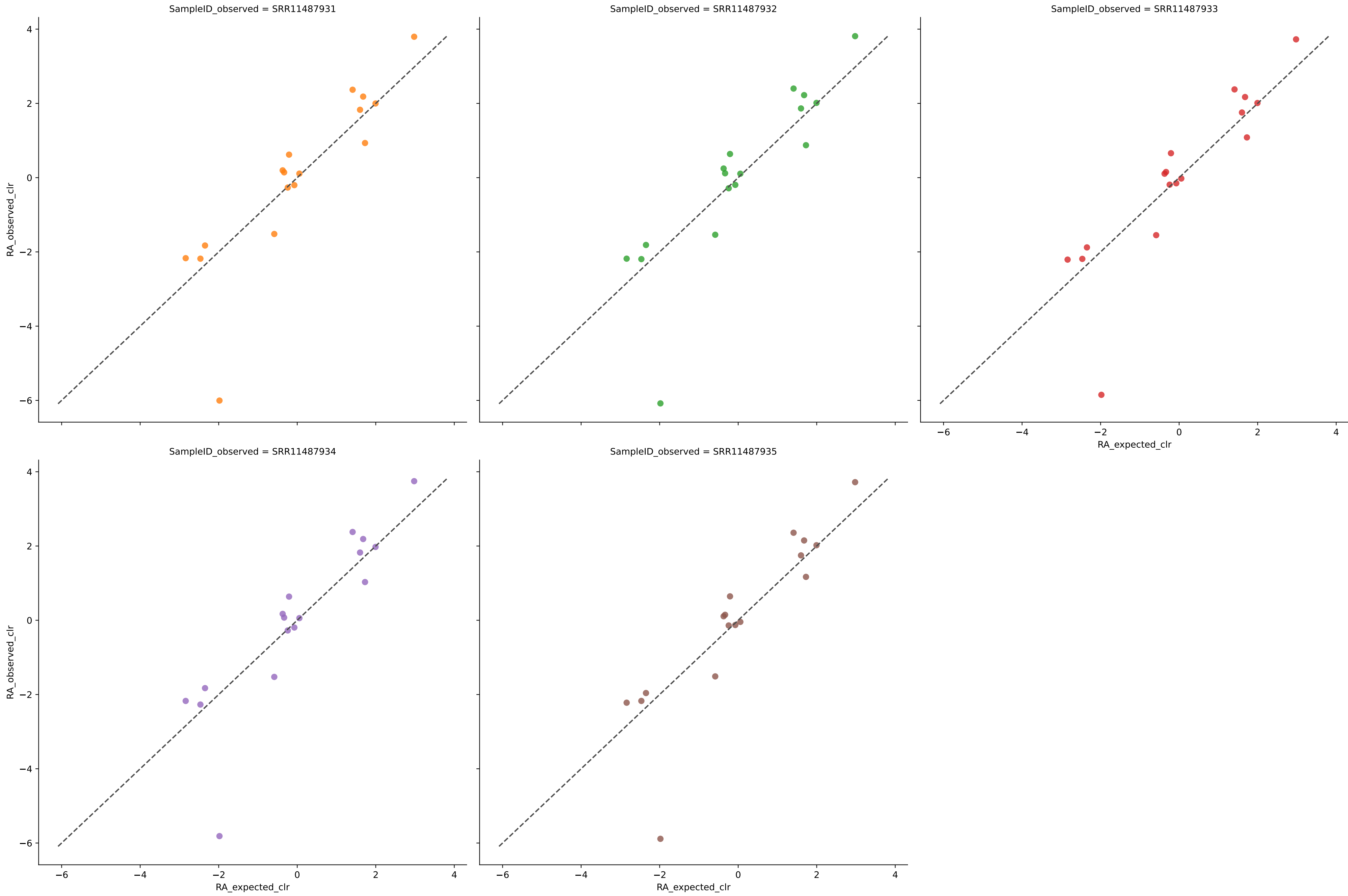
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487931	17	0.9142	0.0034	5.2528	0.8570	0.0060	100.0000	0.0000
SRR11487932	17	0.9028	0.0033	4.8354	0.8613	0.0062	100.0000	0.0000
SRR11487933	17	0.9066	0.0030	6.2762	0.8717	0.0058	100.0000	0.0000
SRR11487934	17	0.8944	0.0032	5.8814	0.8635	0.0062	100.0000	0.0000
SRR11487935	17	0.9113	0.0029	5.1244	0.8775	0.0056	100.0000	0.0000
Average	17	0.9059	0.0031	5.4740	0.8662	0.0060	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using jams in Experiment Amos hilo with filter 0



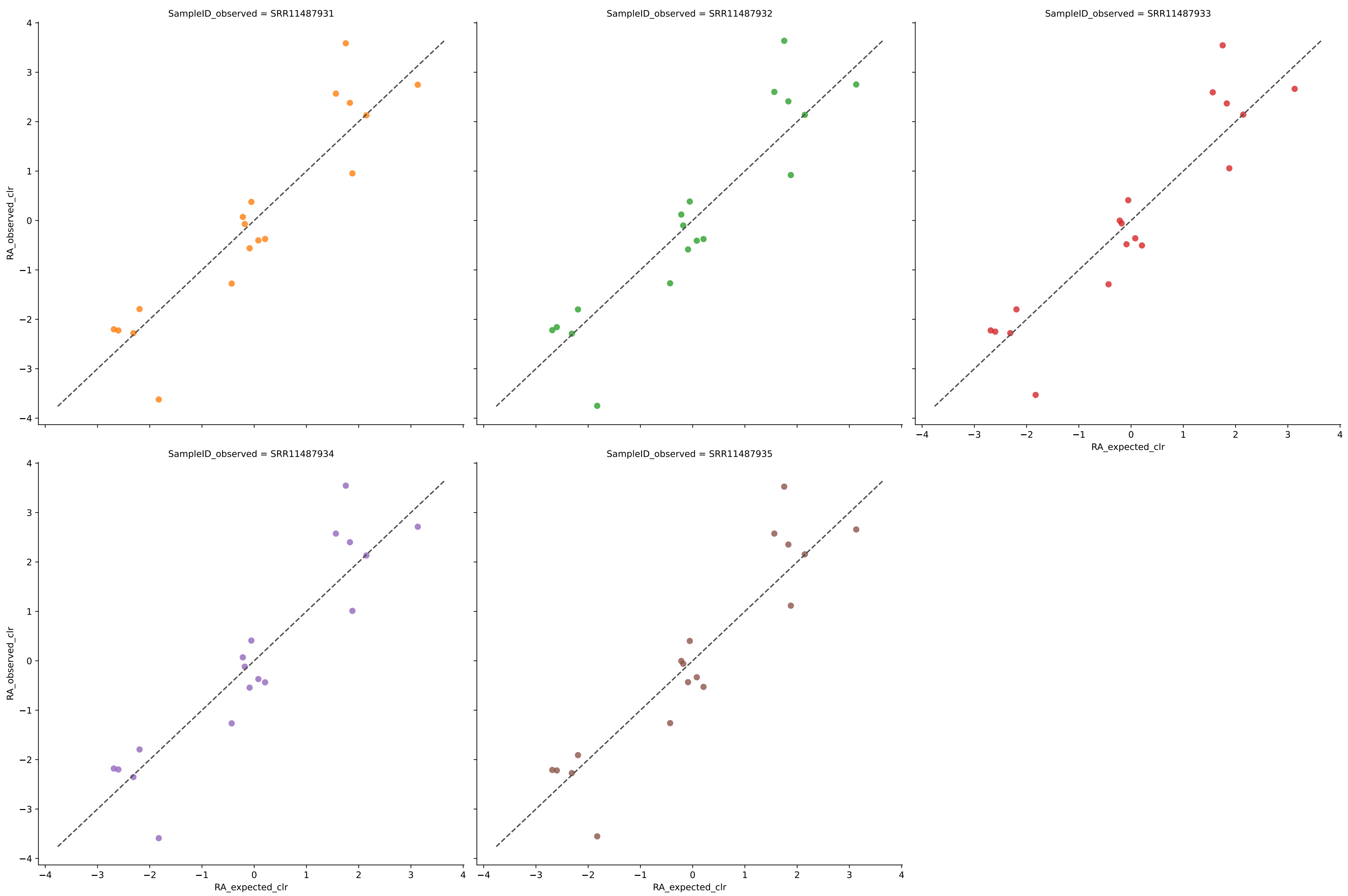
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487931	19	0.9021	0.0041	9.1725	0.8072	0.0067	100.0000	0.0000
SRR11487932	19	0.8961	0.0041	9.5318	0.8047	0.0068	100.0000	0.0000
SRR11487933	19	0.9069	0.0039	8.8944	0.8143	0.0063	100.0000	0.0000
SRR11487934	18	0.8980	0.0042	7.3804	0.8096	0.0067	100.0000	0.0000
SRR11487935	19	0.9133	0.0038	8.8004	0.8172	0.0062	100.0000	0.0000
Average	19	0.9033	0.0040	8.7559	0.8106	0.0065	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using wgsa in Experiment Amos hilo with filter 0



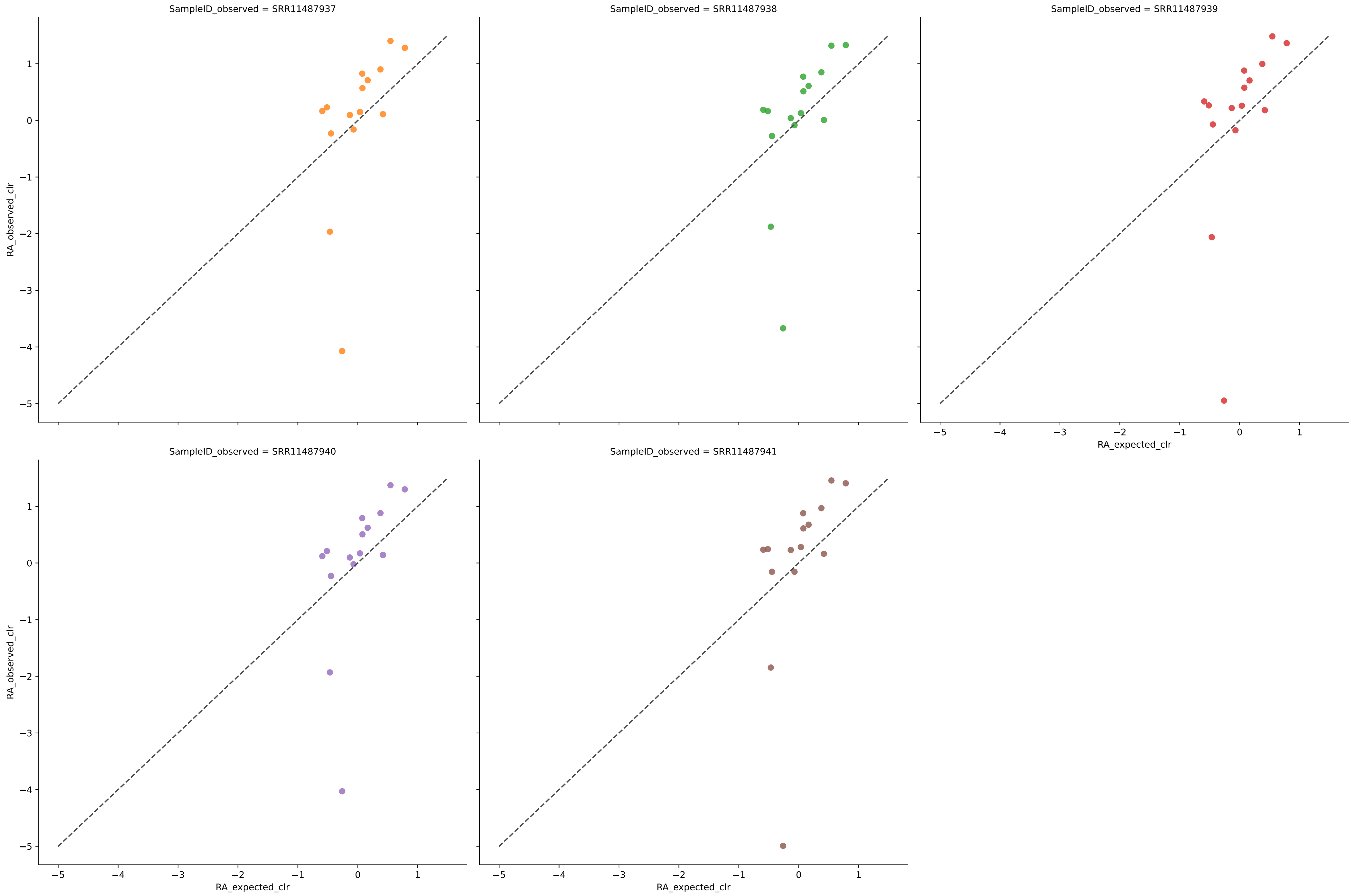
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487931	17	0.9122	0.0044	4.6517	0.8125	0.0084	100.0000	0.0000
SRR11487932	17	0.9108	0.0044	4.7533	0.8110	0.0083	100.0000	0.0000
SRR11487933	17	0.9187	0.0042	4.4721	0.8210	0.0077	100.0000	0.0000
SRR11487934	17	0.9158	0.0043	4.4548	0.8173	0.0079	100.0000	0.0000
SRR11487935	17	0.9223	0.0041	4.4669	0.8255	0.0076	100.0000	0.0000
Average	17	0.9160	0.0043	4.5598	0.8175	0.0080	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using wol in Experiment Amos hilo with filter 0



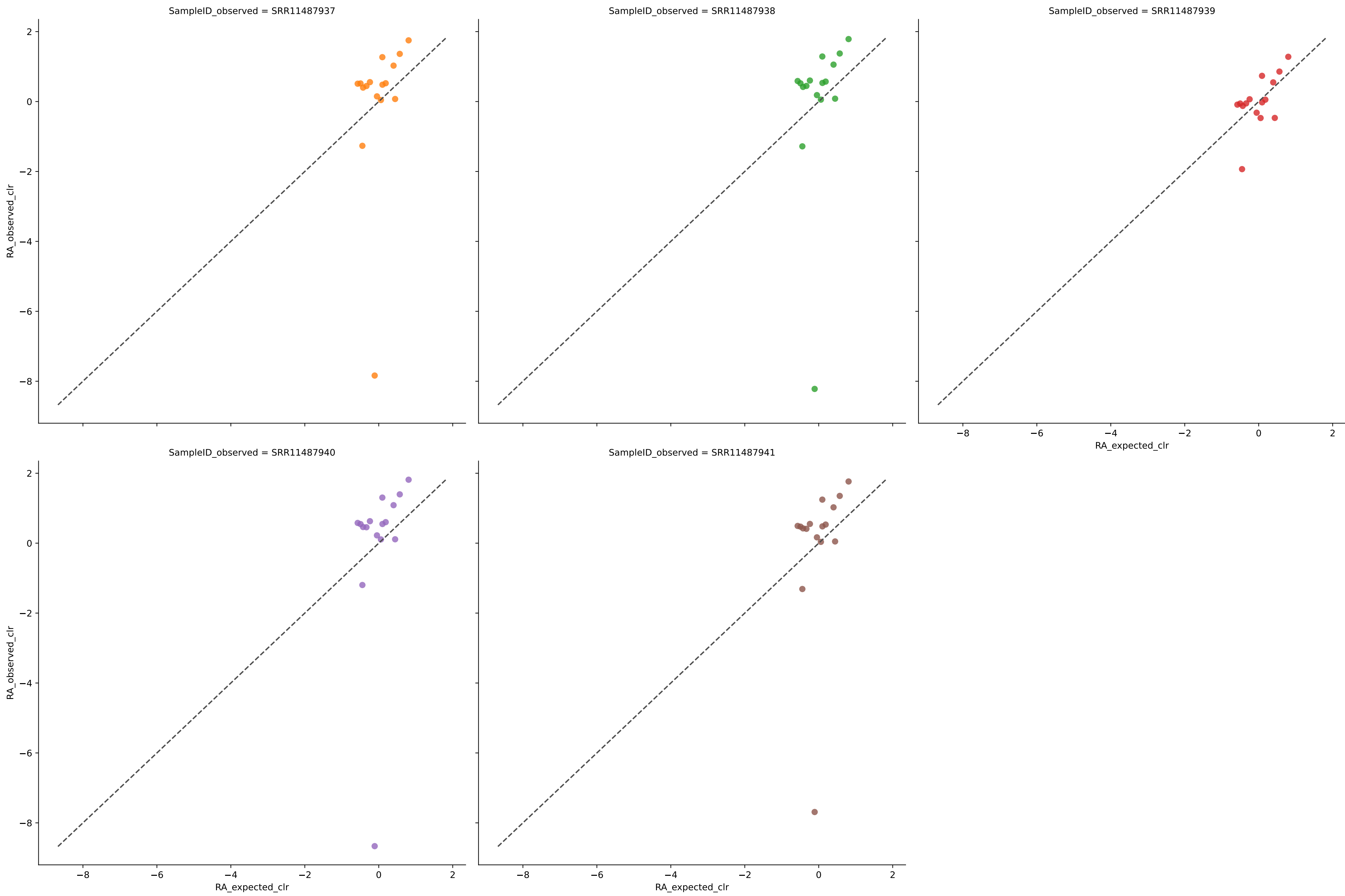
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487931	18	0.2914	0.0083	3.3507	0.6248	0.0177	100.0000	0.0000
SRR11487932	18	0.2756	0.0085	3.4846	0.6160	0.0181	100.0000	0.0000
SRR11487933	18	0.2816	0.0084	3.2745	0.6226	0.0177	100.0000	0.0000
SRR11487934	18	0.2963	0.0083	3.3150	0.6261	0.0175	100.0000	0.0000
SRR11487935	18	0.2883	0.0083	3.2343	0.6285	0.0176	100.0000	0.0000
Average	18	0.2866	0.0084	3.3318	0.6236	0.0177	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using bio4 in Experiment Amos mixed with filter 0



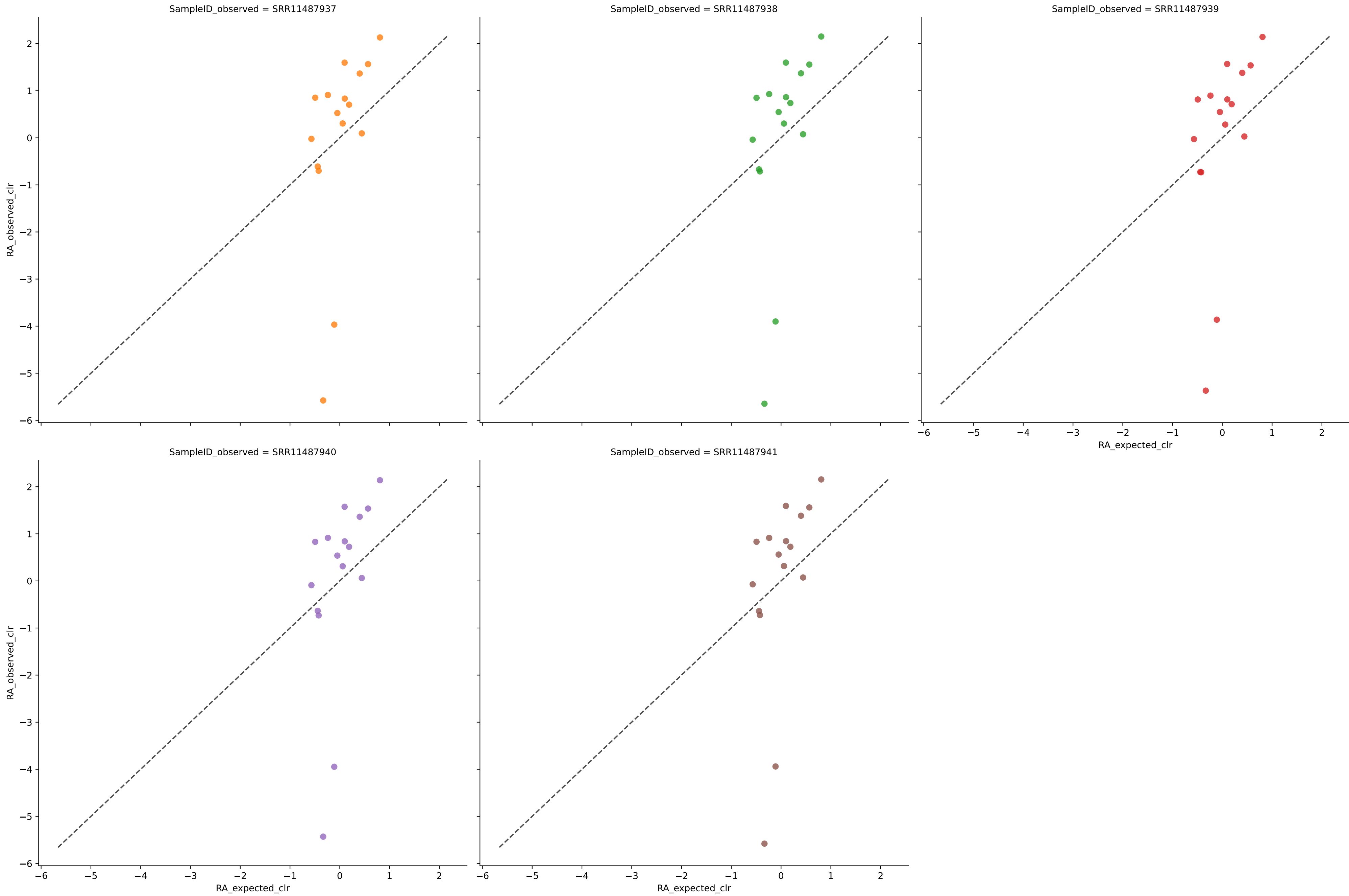
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487937	15	0.6595	0.0047	4.5249	0.8236	0.0058	100.0000	0.0000
SRR11487938	15	0.6856	0.0046	4.1118	0.8260	0.0056	100.0000	0.0000
SRR11487939	15	0.6543	0.0046	5.3975	0.8269	0.0058	100.0000	0.0000
SRR11487940	15	0.6903	0.0044	4.4374	0.8352	0.0055	100.0000	0.0000
SRR11487941	15	0.6856	0.0046	5.3530	0.8283	0.0057	100.0000	0.0000
Average	15	0.6751	0.0046	4.7649	0.8280	0.0057	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using jams in Experiment Amos mixed with filter 0



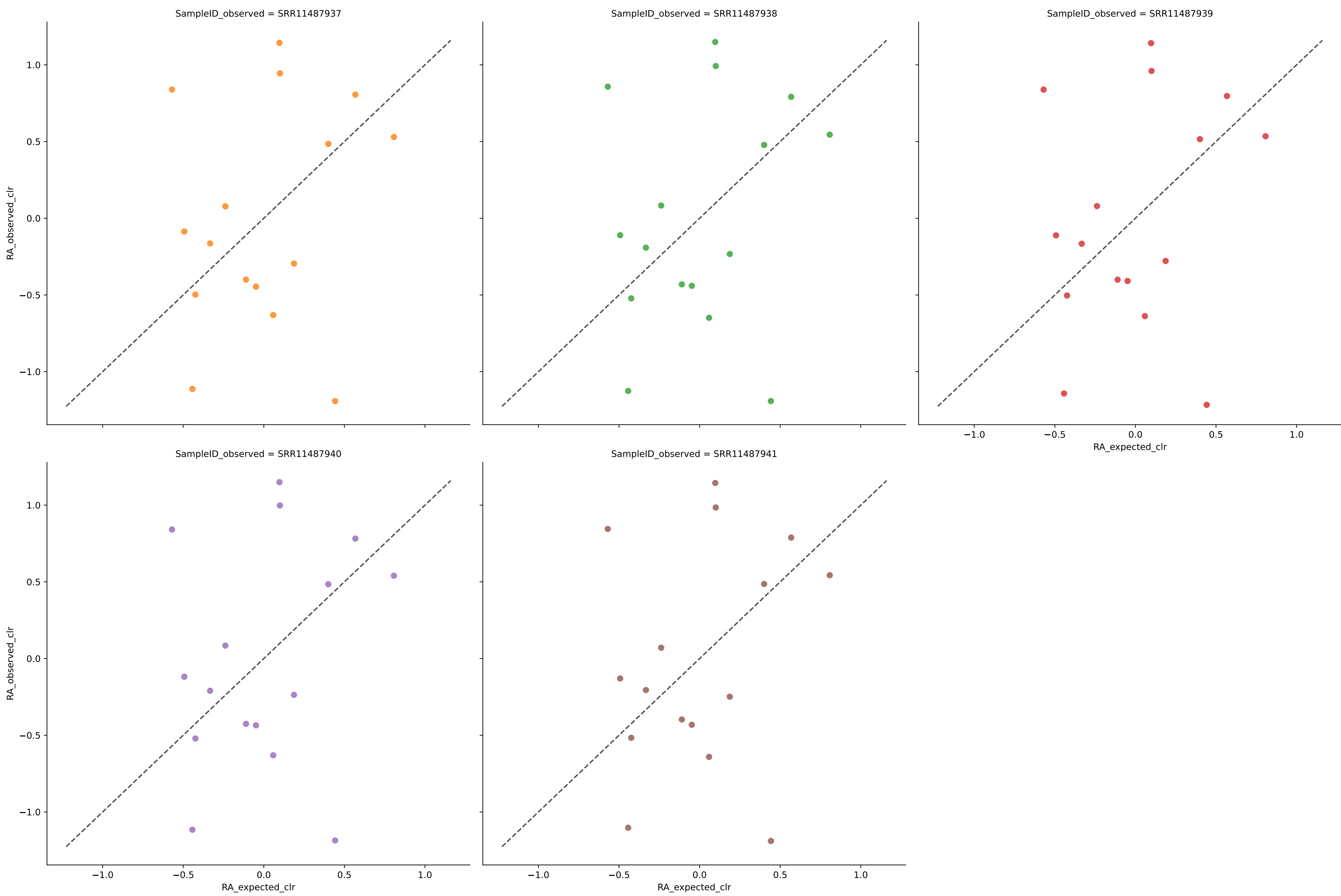
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487937	16	0.5479	0.0051	8.2583	0.7959	0.0062	100.0000	0.0000
SRR11487938	16	0.5460	0.0051	8.6504	0.7958	0.0062	100.0000	0.0000
SRR11487939	15	0.5984	0.0047	2.2036	0.8197	0.0058	100.0000	0.0000
SRR11487940	16	0.5531	0.0051	9.0831	0.7977	0.0062	100.0000	0.0000
SRR11487941	16	0.5581	0.0051	8.1152	0.7969	0.0062	100.0000	0.0000
Average	16	0.5607	0.0050	7.2621	0.8012	0.0061	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using wgsa in Experiment Amos mixed with filter 0



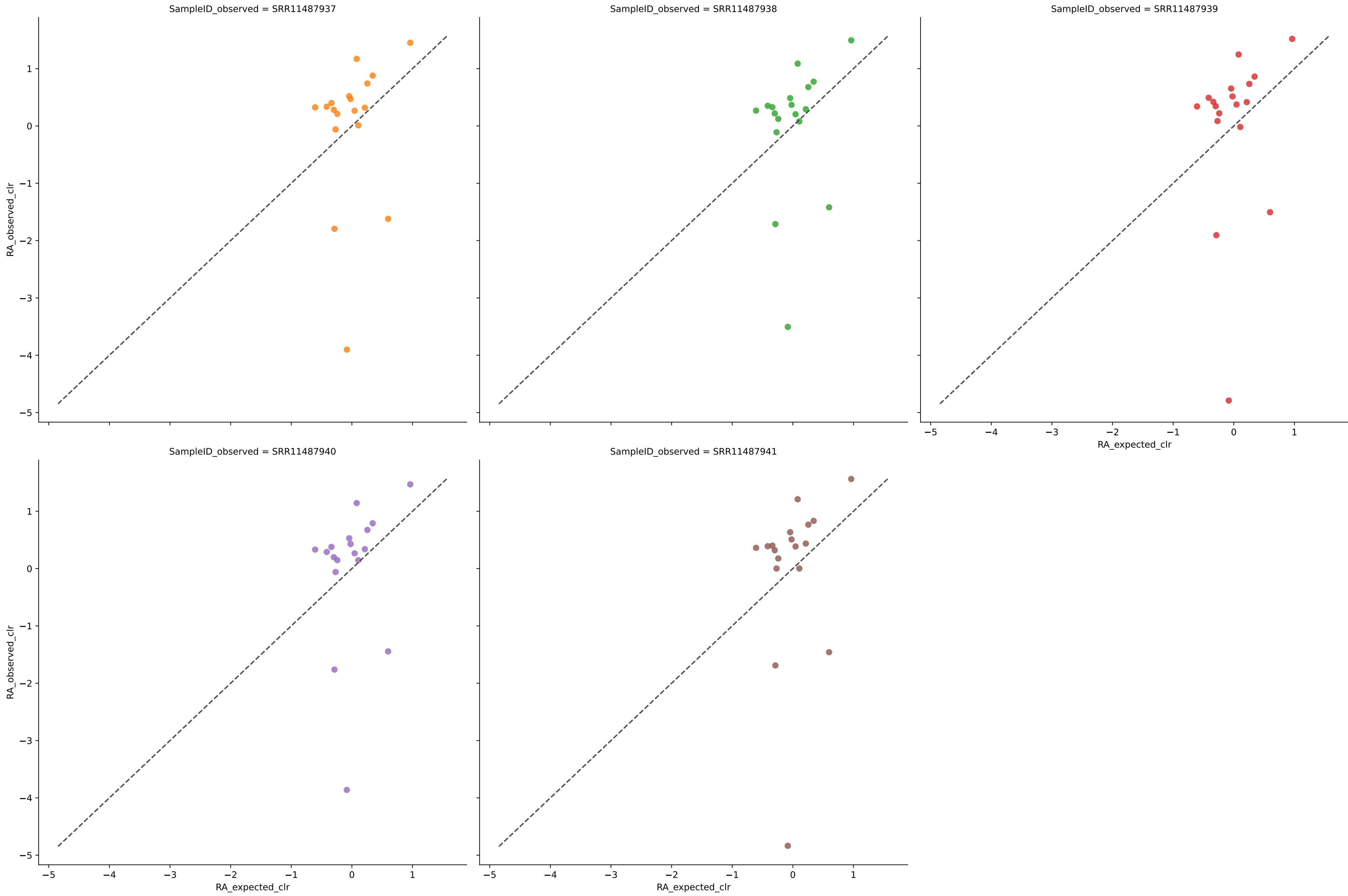
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487937	16	0.6162	0.0064	7.2906	0.7422	0.0081	100.0000	0.0000
SRR11487938	16	0.6172	0.0064	7.3210	0.7424	0.0082	100.0000	0.0000
SRR11487939	16	0.6217	0.0065	7.0792	0.7405	0.0082	100.0000	0.0000
SRR11487940	16	0.6200	0.0064	7.1672	0.7428	0.0081	100.0000	0.0000
SRR11487941	16	0.6231	0.0065	7.2852	0.7413	0.0082	100.0000	0.0000
Average	16	0.6196	0.0065	7.2286	0.7418	0.0082	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using wol in Experiment Amos mixed with filter 0



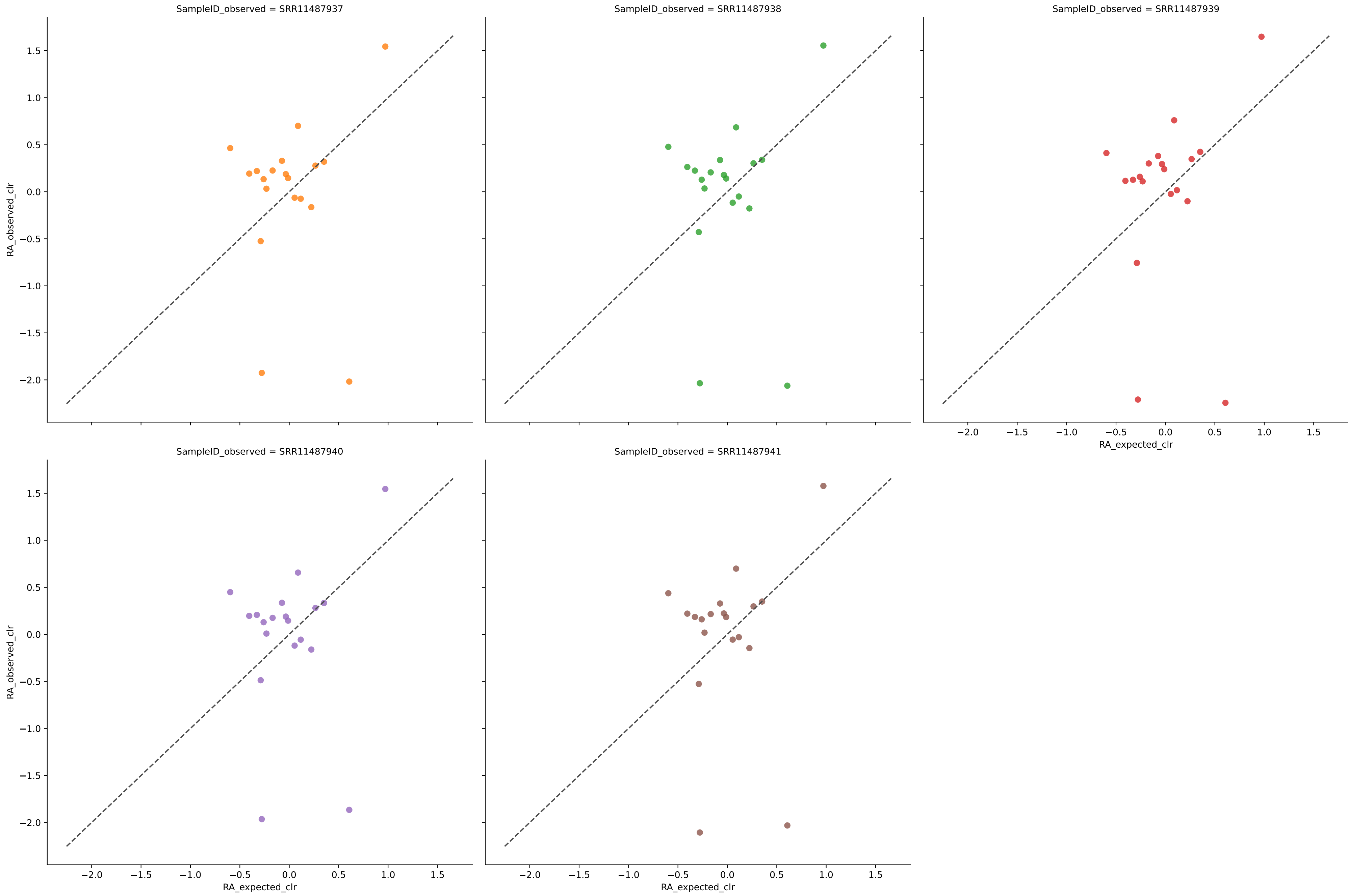
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487937	16	0.0645	0.0066	2.8809	0.7364	0.0087	100.0000	0.0000
SRR11487938	16	0.0633	0.0066	2.9006	0.7340	0.0089	100.0000	0.0000
SRR11487939	16	0.0662	0.0065	2.8965	0.7387	0.0087	100.0000	0.0000
SRR11487940	16	0.0643	0.0066	2.8819	0.7351	0.0088	100.0000	0.0000
SRR11487941	16	0.0658	0.0066	2.8748	0.7369	0.0088	100.0000	0.0000
Average	16	0.0648	0.0066	2.8869	0.7362	0.0088	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using bio4 in Experiment Amos mixed with filter 0



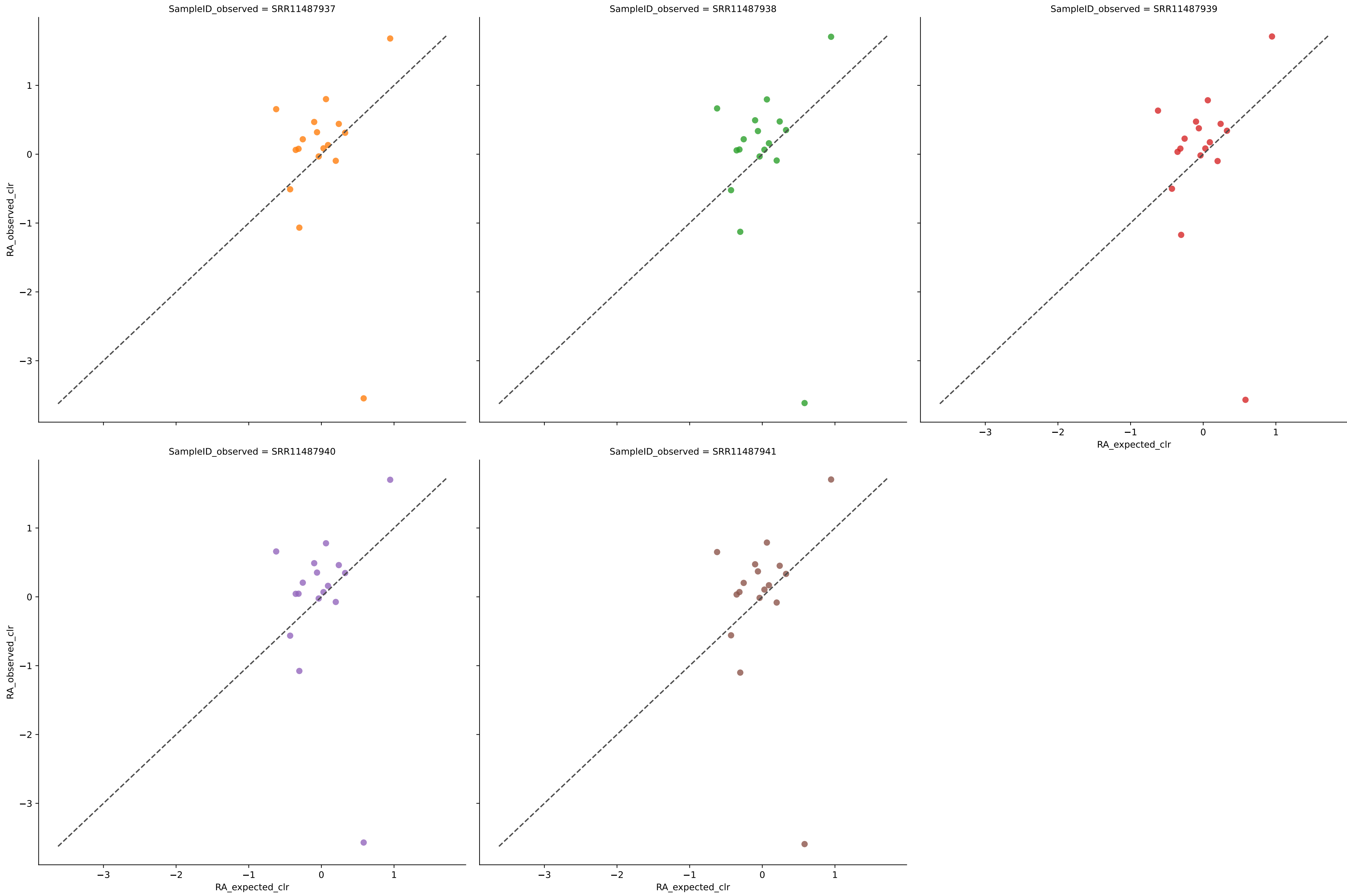
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487937	18	0.2996	0.0046	5.1875	0.7914	0.0063	100.0000	0.0000
SRR11487938	18	0.3547	0.0045	4.7066	0.7969	0.0062	100.0000	0.0000
SRR11487939	18	0.2928	0.0046	5.9450	0.7951	0.0063	100.0000	0.0000
SRR11487940	18	0.3277	0.0044	5.0307	0.8030	0.0061	100.0000	0.0000
SRR11487941	18	0.3332	0.0045	5.8823	0.7971	0.0063	100.0000	0.0000
Average	18	0.3216	0.0045	5.3504	0.7967	0.0062	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using jams in Experiment Amos mixed with filter 0



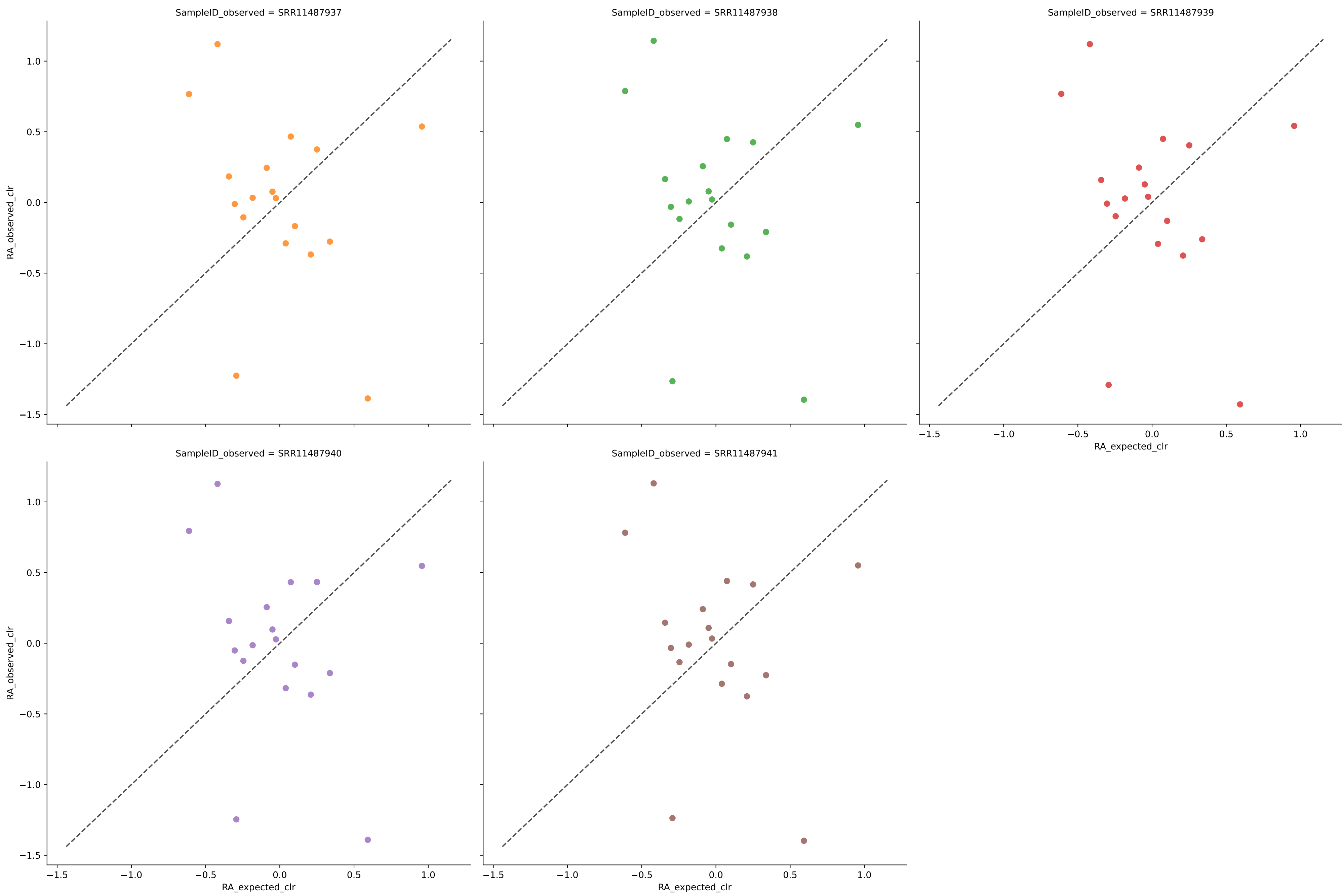
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487937	19	0.3708	0.0043	3.6003	0.7947	0.0060	100.0000	0.0000
SRR11487938	19	0.3682	0.0044	3.6950	0.7941	0.0061	100.0000	0.0000
SRR11487939	19	0.4162	0.0042	3.9369	0.7974	0.0060	100.0000	0.0000
SRR11487940	19	0.3834	0.0043	3.4885	0.7968	0.0060	100.0000	0.0000
SRR11487941	19	0.3887	0.0043	3.6944	0.7964	0.0060	100.0000	0.0000
Average	19	0.3854	0.0043	3.6830	0.7959	0.0060	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using wgsa in Experiment Amos mixed with filter 0



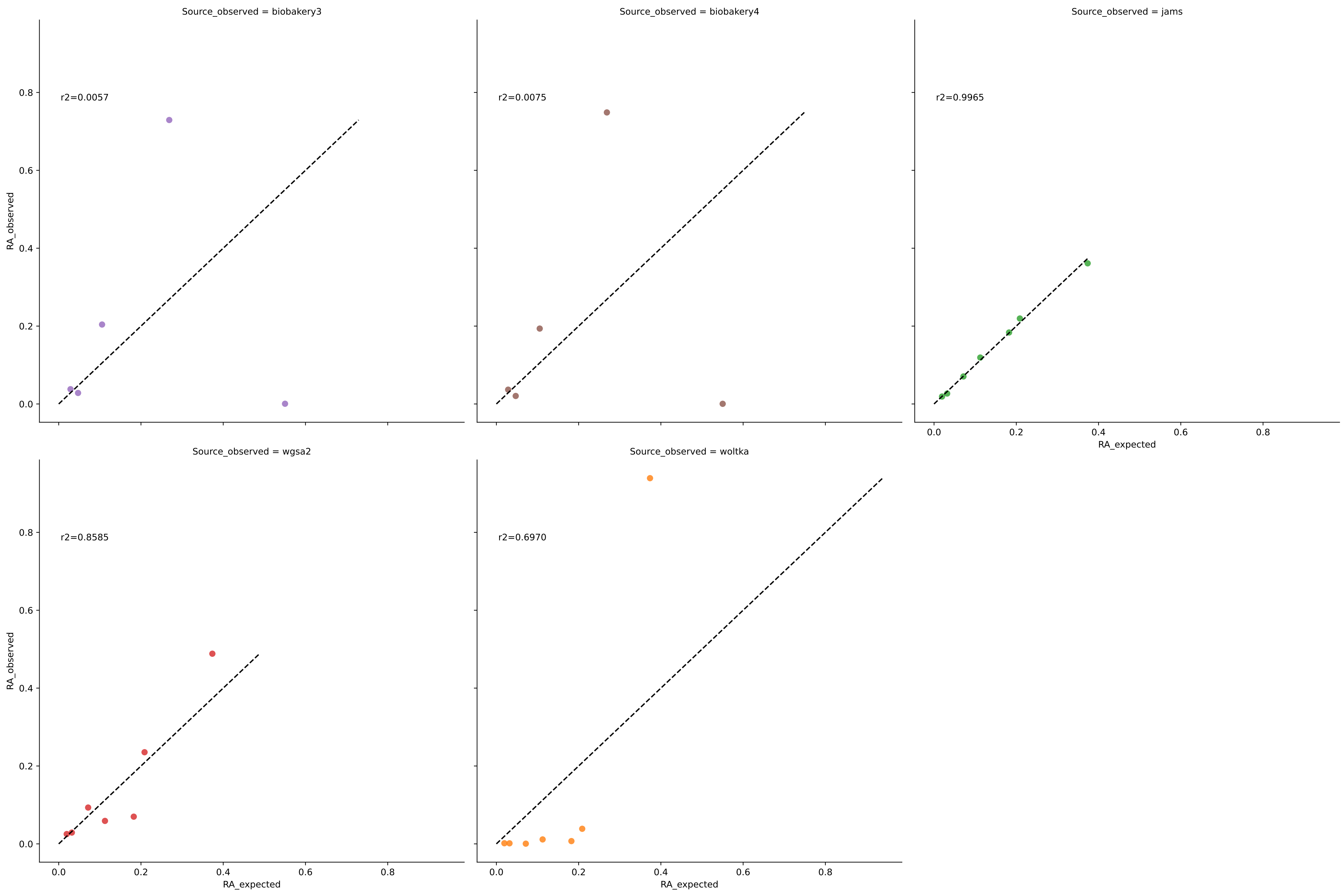
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487937	17	0.3833	0.0051	4.6362	0.7831	0.0074	100.0000	0.0000
SRR11487938	17	0.3924	0.0051	4.7186	0.7829	0.0074	100.0000	0.0000
SRR11487939	17	0.3983	0.0051	4.6751	0.7827	0.0075	100.0000	0.0000
SRR11487940	17	0.3953	0.0051	4.6599	0.7843	0.0074	100.0000	0.0000
SRR11487941	17	0.3961	0.0051	4.6848	0.7841	0.0074	100.0000	0.0000
Average	17	0.3931	0.0051	4.6749	0.7834	0.0074	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using wol in Experiment Amos mixed with filter 0

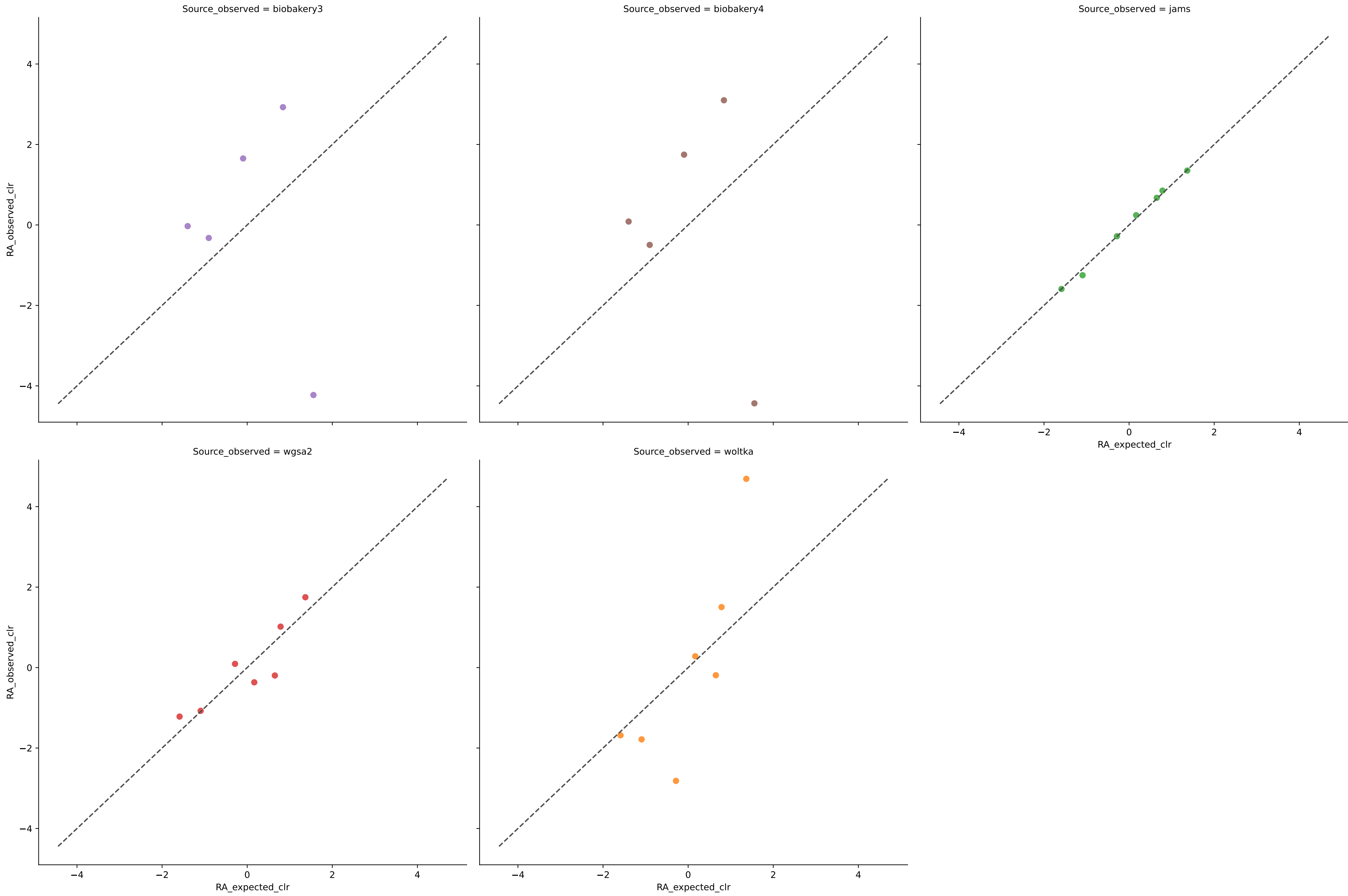


	Diversity	R ²	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487937	18	0.0234	0.0058	3.2948	0.7376	0.0085	100.0000	0.0000
SRR11487938	18	0.0204	0.0058	3.3182	0.7373	0.0085	100.0000	0.0000
SRR11487939	18	0.0221	0.0058	3.3334	0.7377	0.0084	100.0000	0.0000
SRR11487940	18	0.0196	0.0058	3.2964	0.7397	0.0084	100.0000	0.0000
SRR11487941	18	0.0193	0.0058	3.2935	0.7407	0.0084	100.0000	0.0000
Average	18	0.0210	0.0058	3.3073	0.7386	0.0085	100.0000	0.0000

Bivariate Linear Regression for Sample S1 in Experiment bmock12 (Genus at filter threshold 1e-05)

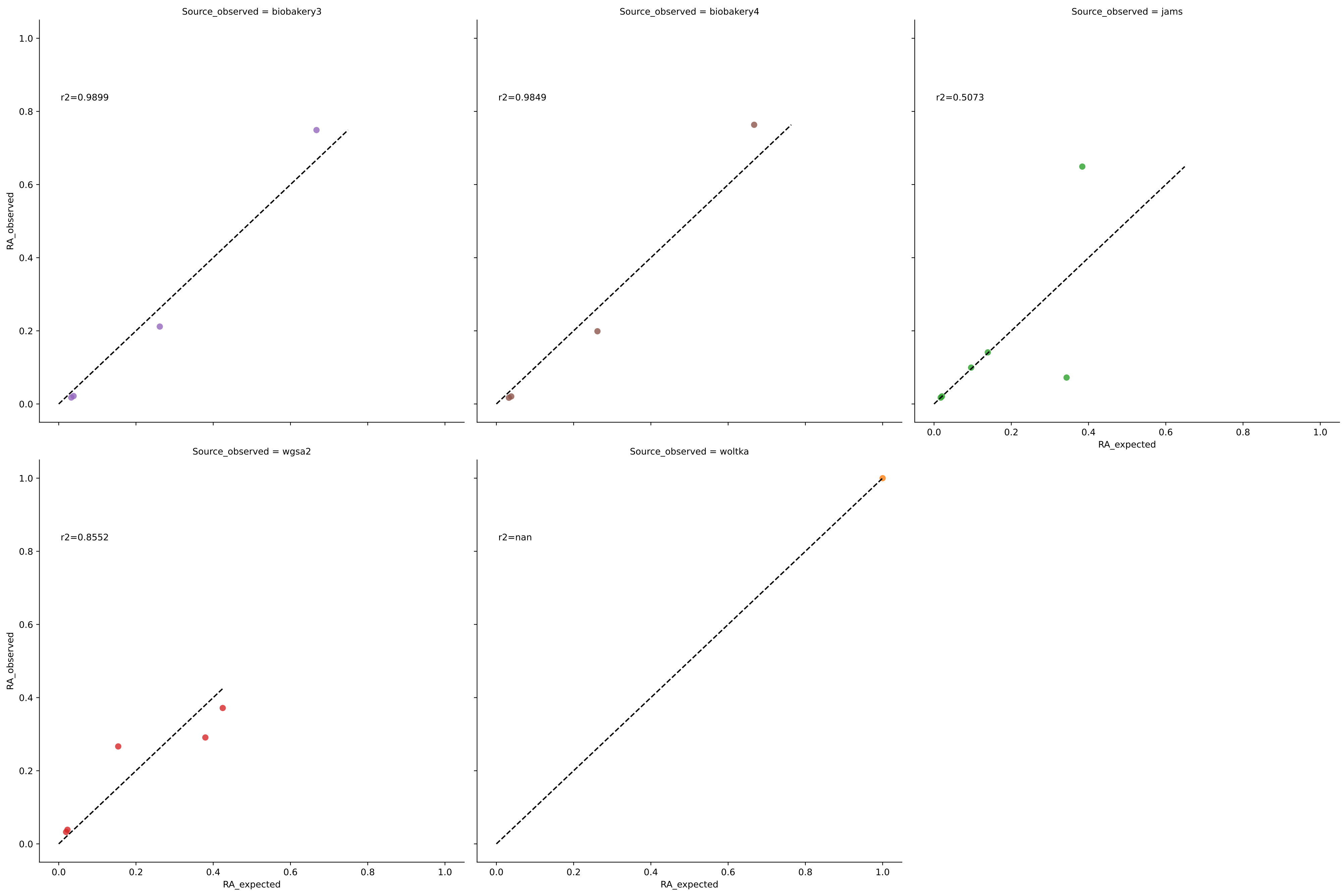


Bivariate Linear Regression for Sample S1 in Experiment bmock12 (Genus at filter threshold 1e-05)

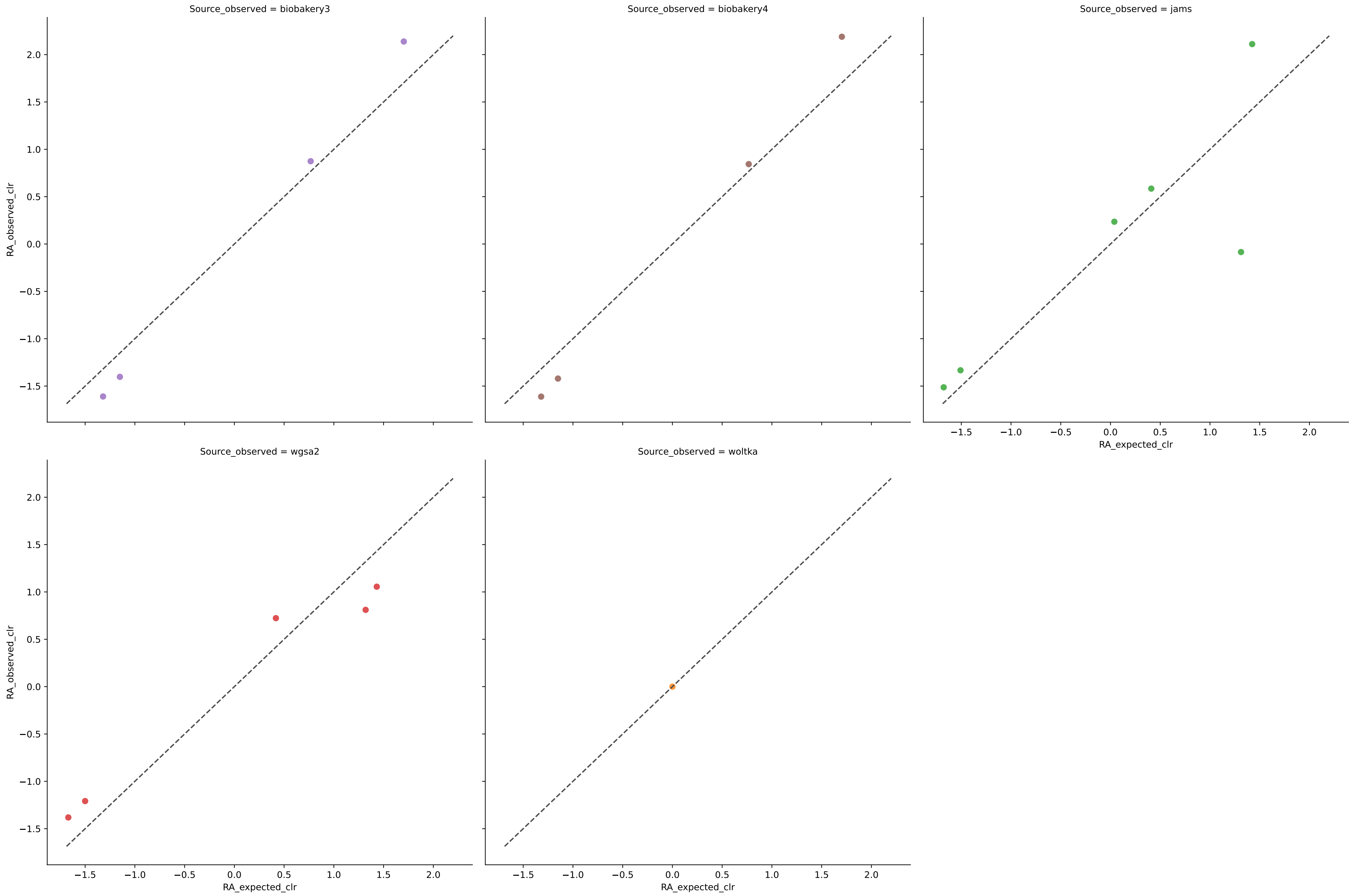


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
biobakery3	5	0.0057	0.2274	6.5633	0.4316	0.3238	100.0000	0.0000
biobakery4	5	0.0075	0.2305	6.8378	0.4237	0.3290	100.0000	0.0000
jams	7	0.9965	0.0053	0.1890	0.9813	0.0070	100.0000	0.0000
wgsa2	7	0.8585	0.0483	1.2171	0.8311	0.0654	100.0000	0.0000
woltka	7	0.6970	0.1616	4.3790	0.4344	0.2379	100.0000	0.0000

Bivariate Linear Regression for Sample S1 in Experiment bmock12 (Species at filter threshold 1e-05)

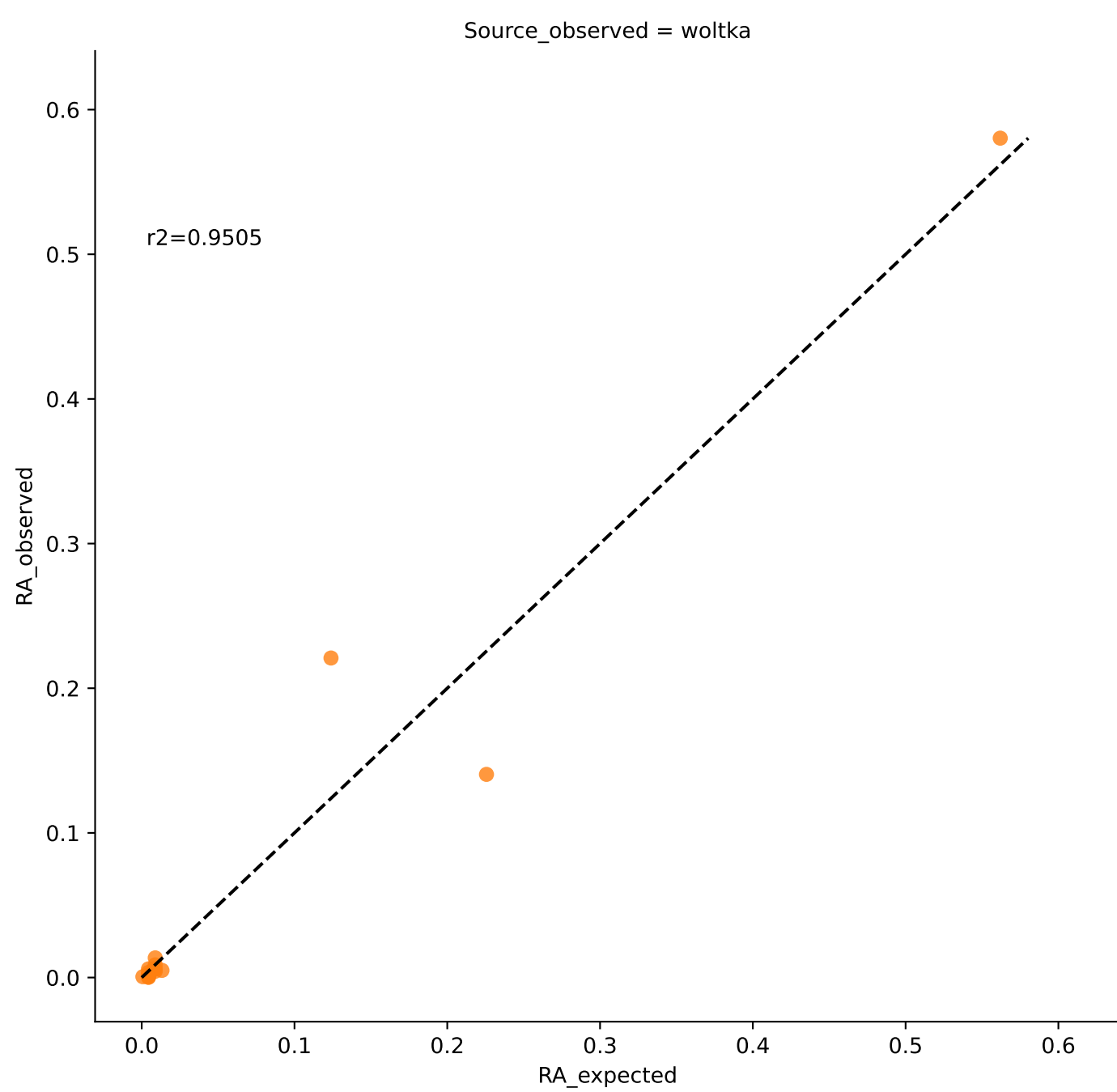
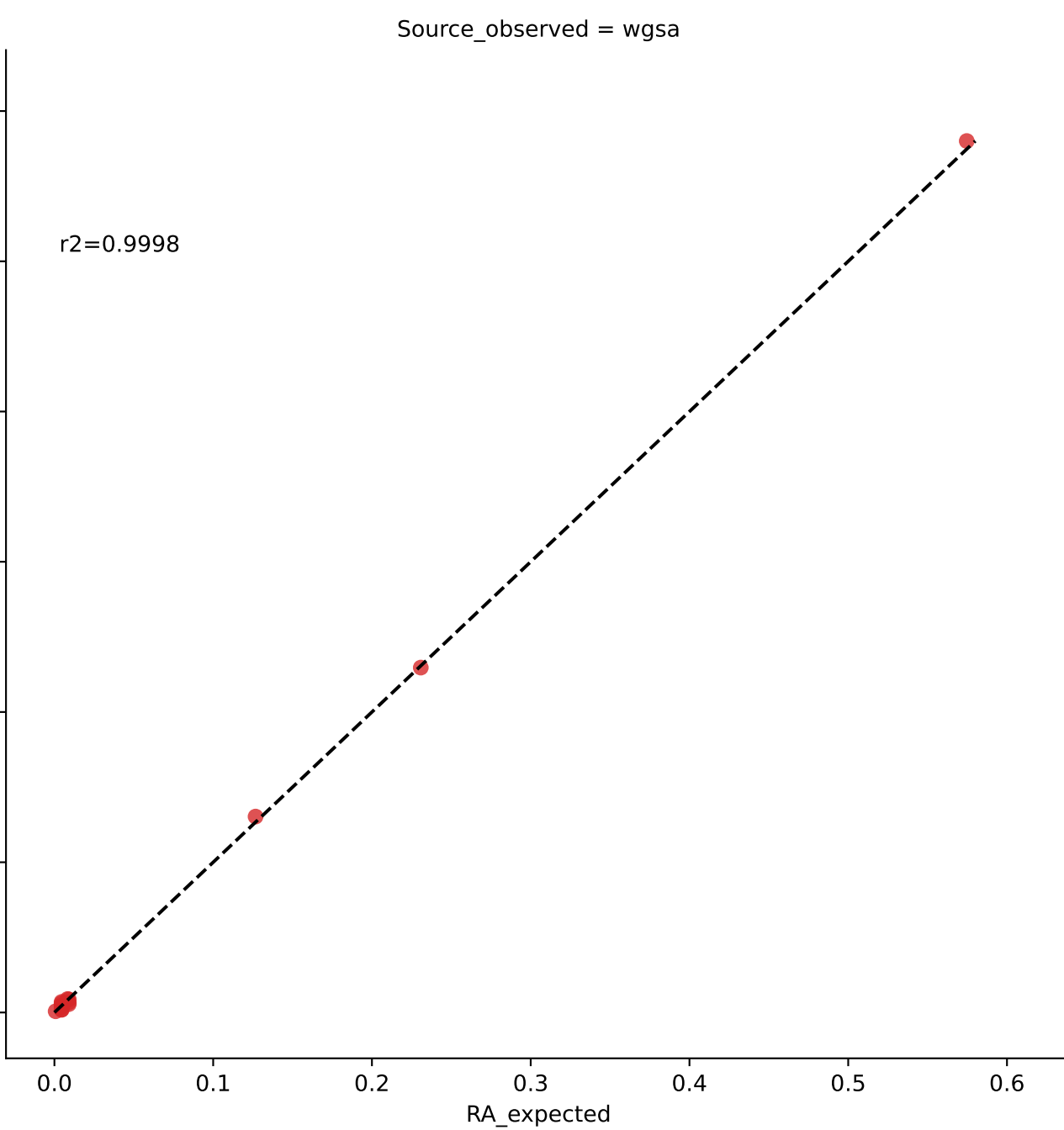
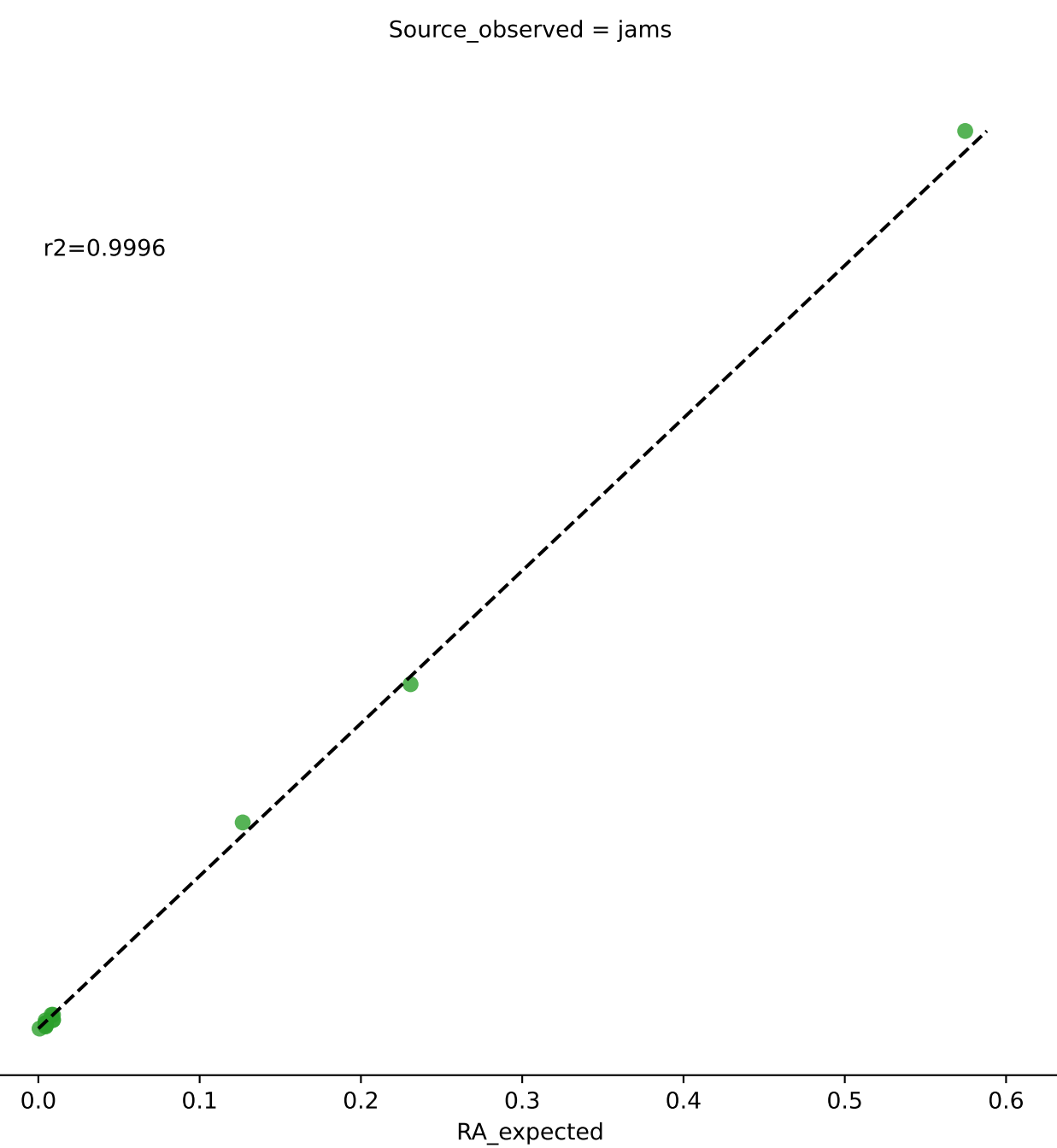
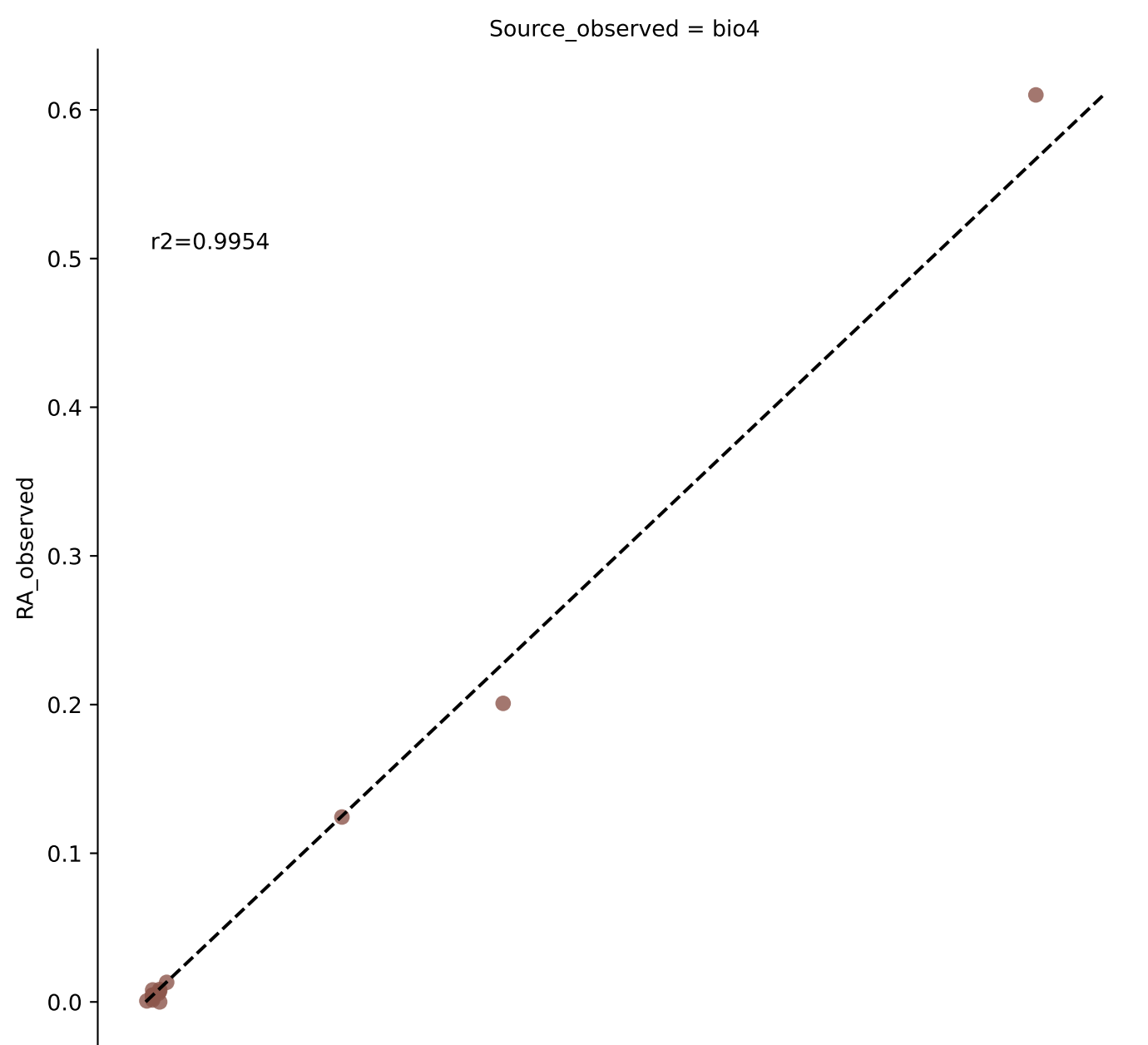


Bivariate Linear Regression for Sample S1 in Experiment bmock12 (Species at filter threshold 1e-05)

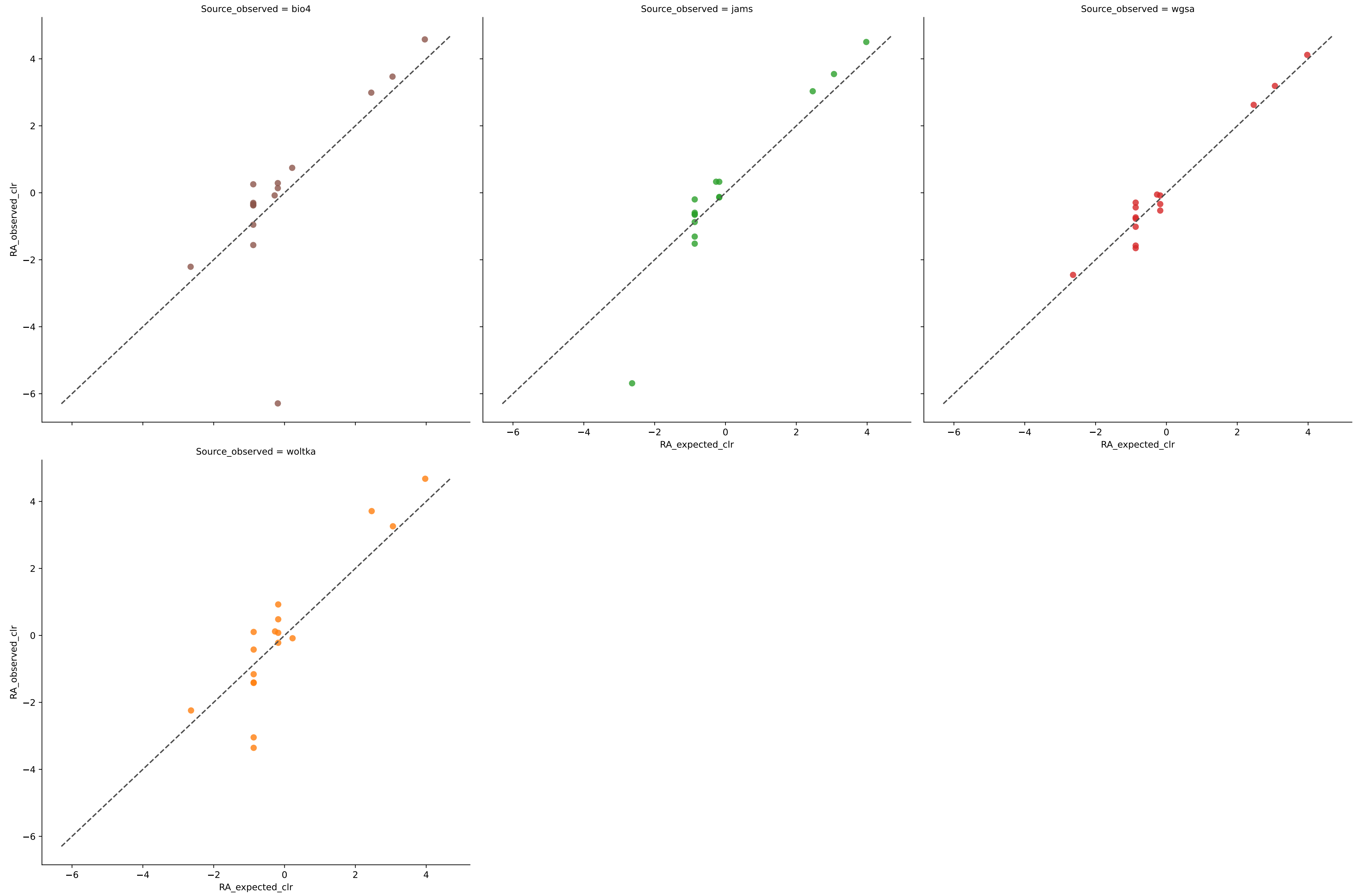


	Diversity	R ²	MAE	AD	1-BC	RMSE	Sens	FPRA
biobakery3	4	0.9899	0.0408	0.5907	0.9184	0.0492	100.0000	0.0000
biobakery4	4	0.9849	0.0480	0.6324	0.9040	0.0586	100.0000	0.0000
jams	6	0.5073	0.0903	1.5970	0.7290	0.1548	100.0000	0.0000
wgsa2	5	0.8552	0.0567	0.8135	0.8582	0.0690	100.0000	0.0000
woltka	1	nan	0.0000	0.0000	1.0000	0.0000	100.0000	0.0000

Bivariate Linear Regression for Sample S1 in Experiment camisimGI (Genus at filter threshold 1e-05)

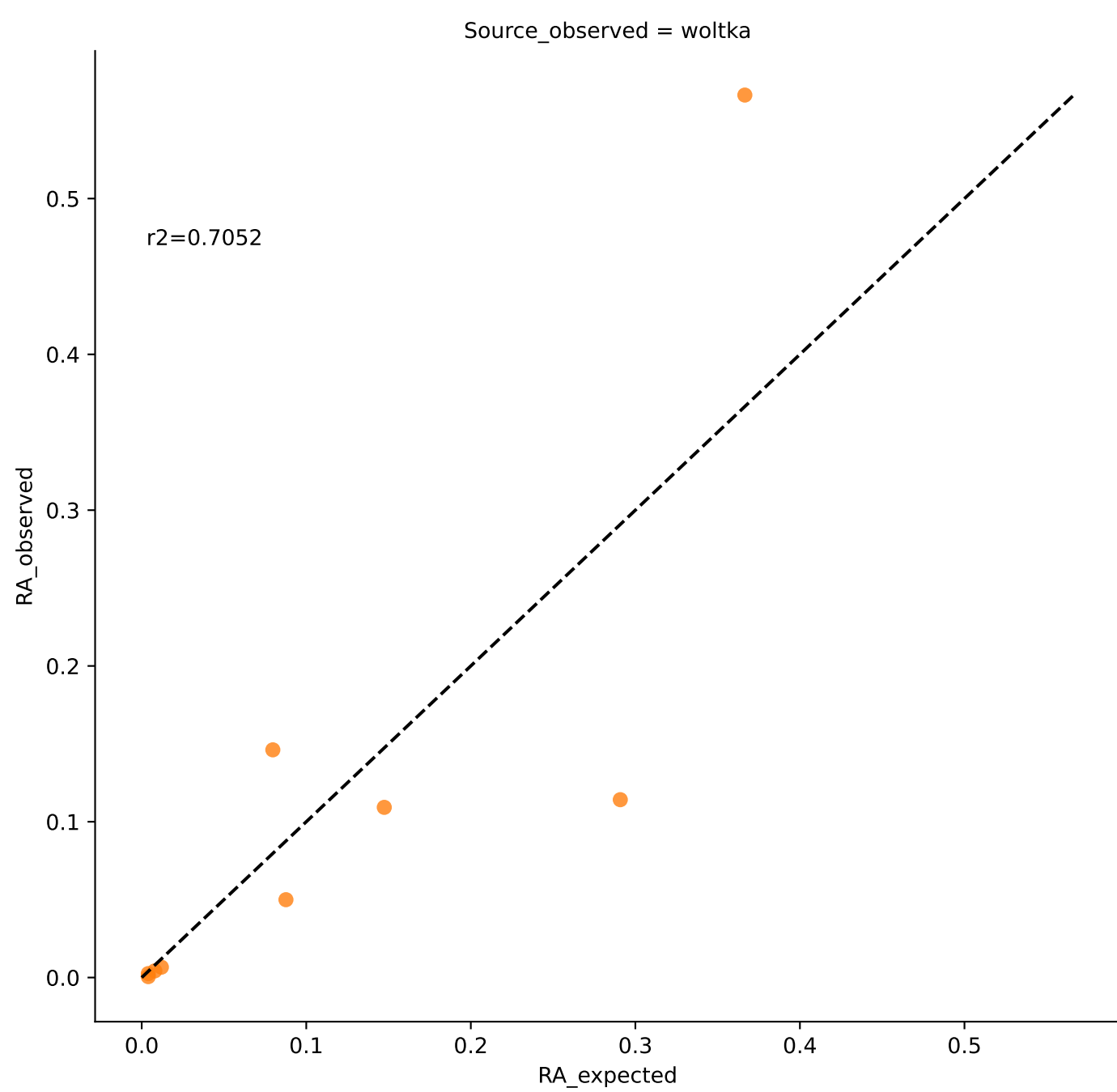
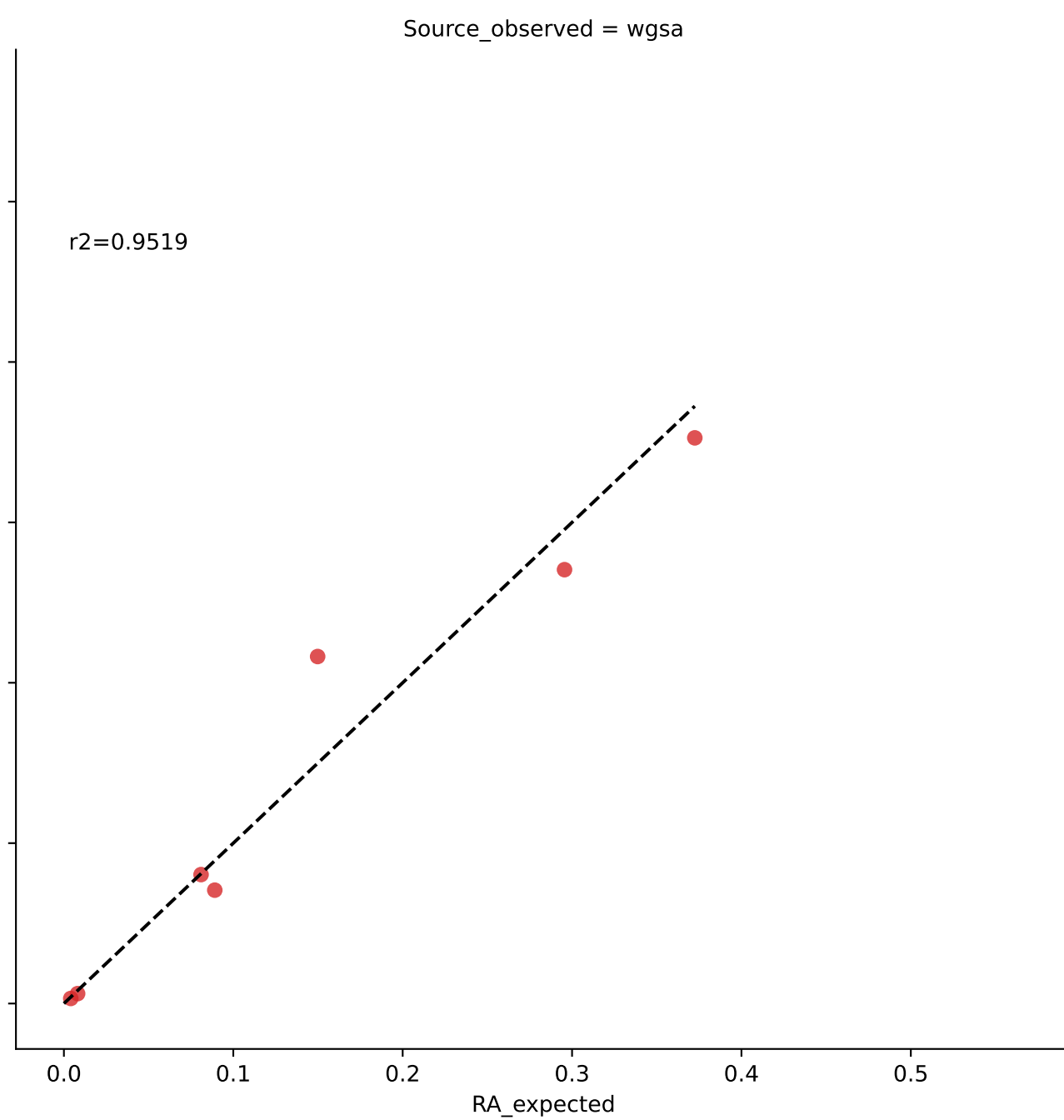
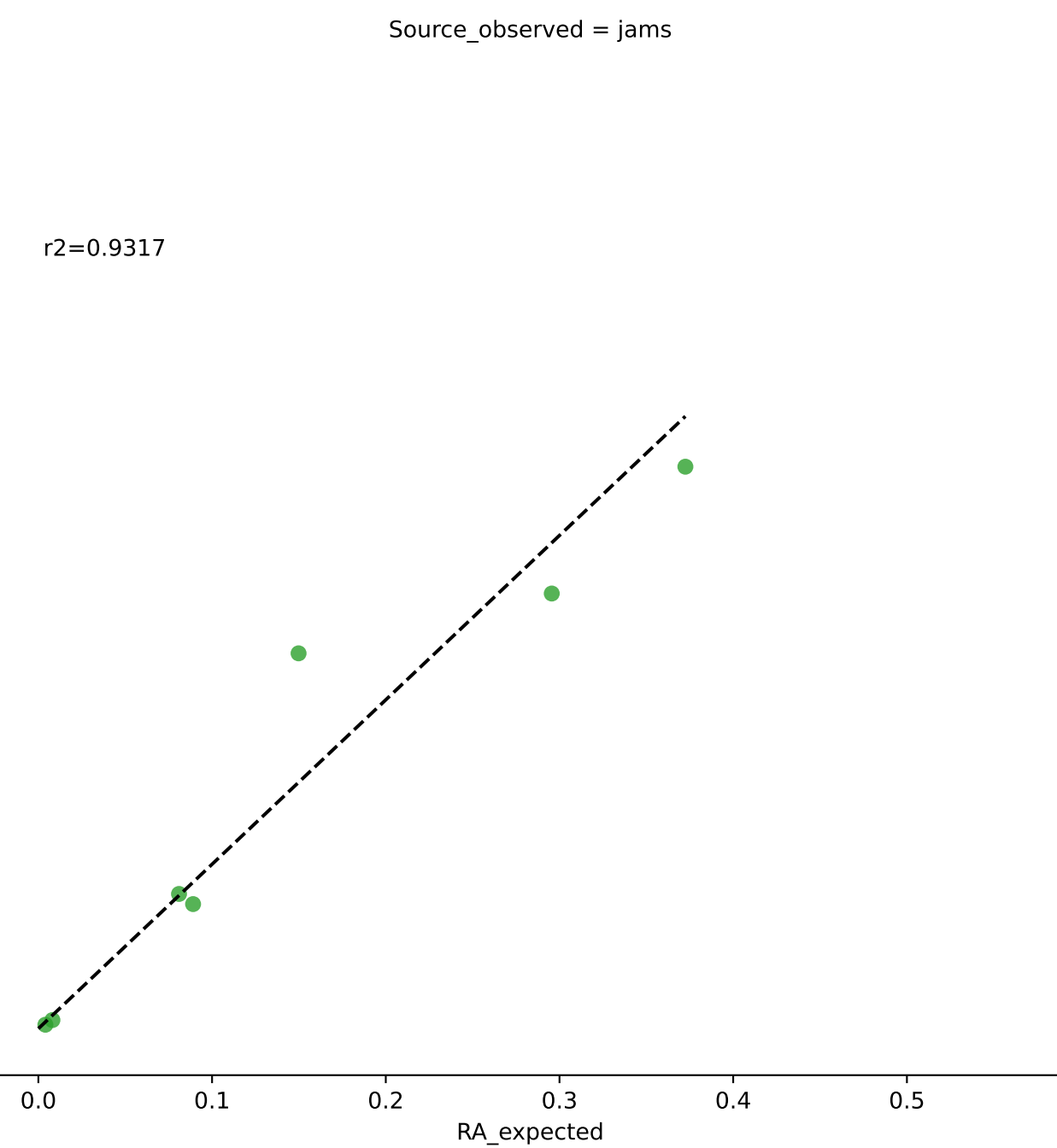
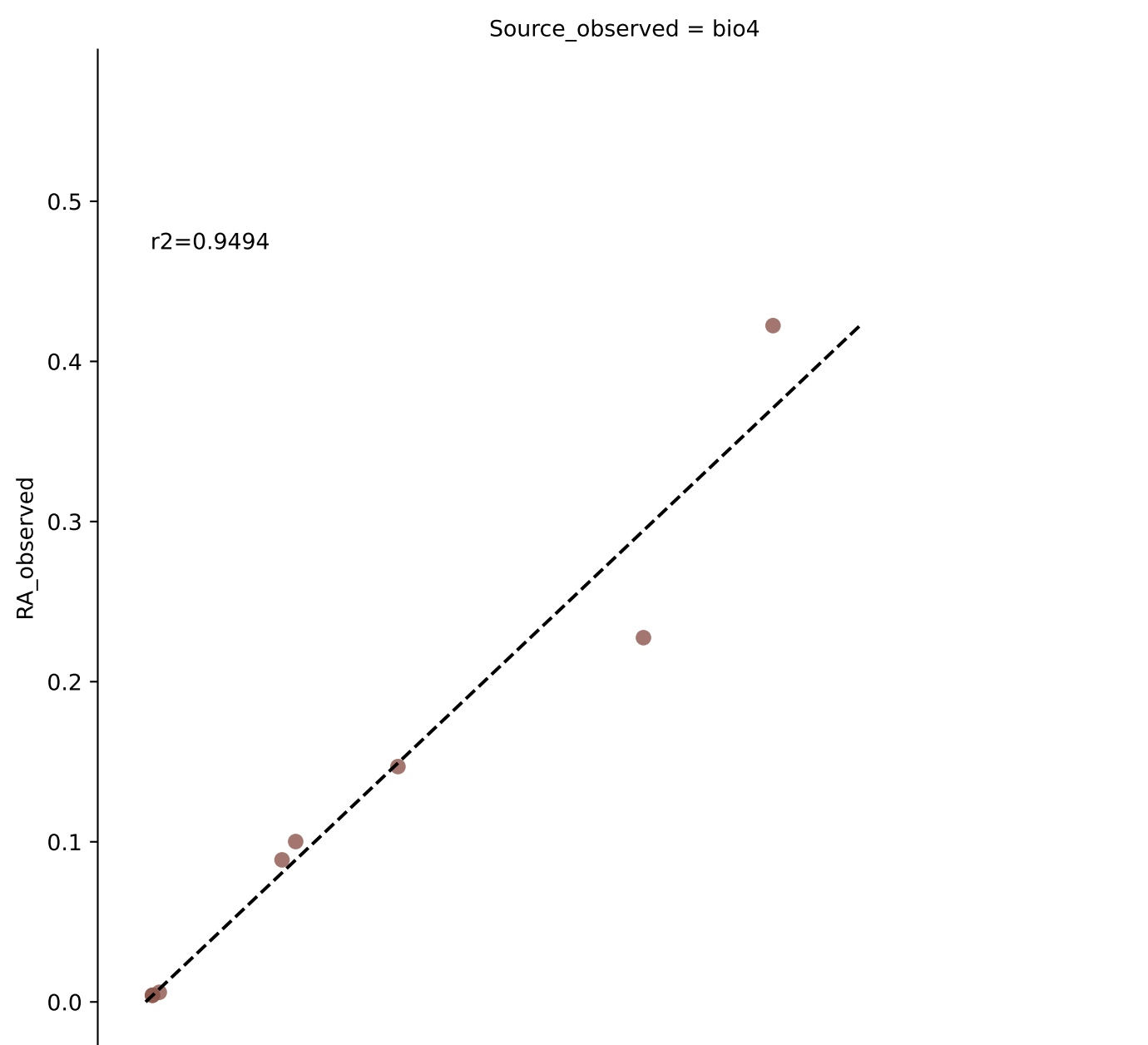


Bivariate Linear Regression for Sample S1 in Experiment camisimGI (Genus at filter threshold 1e-05)

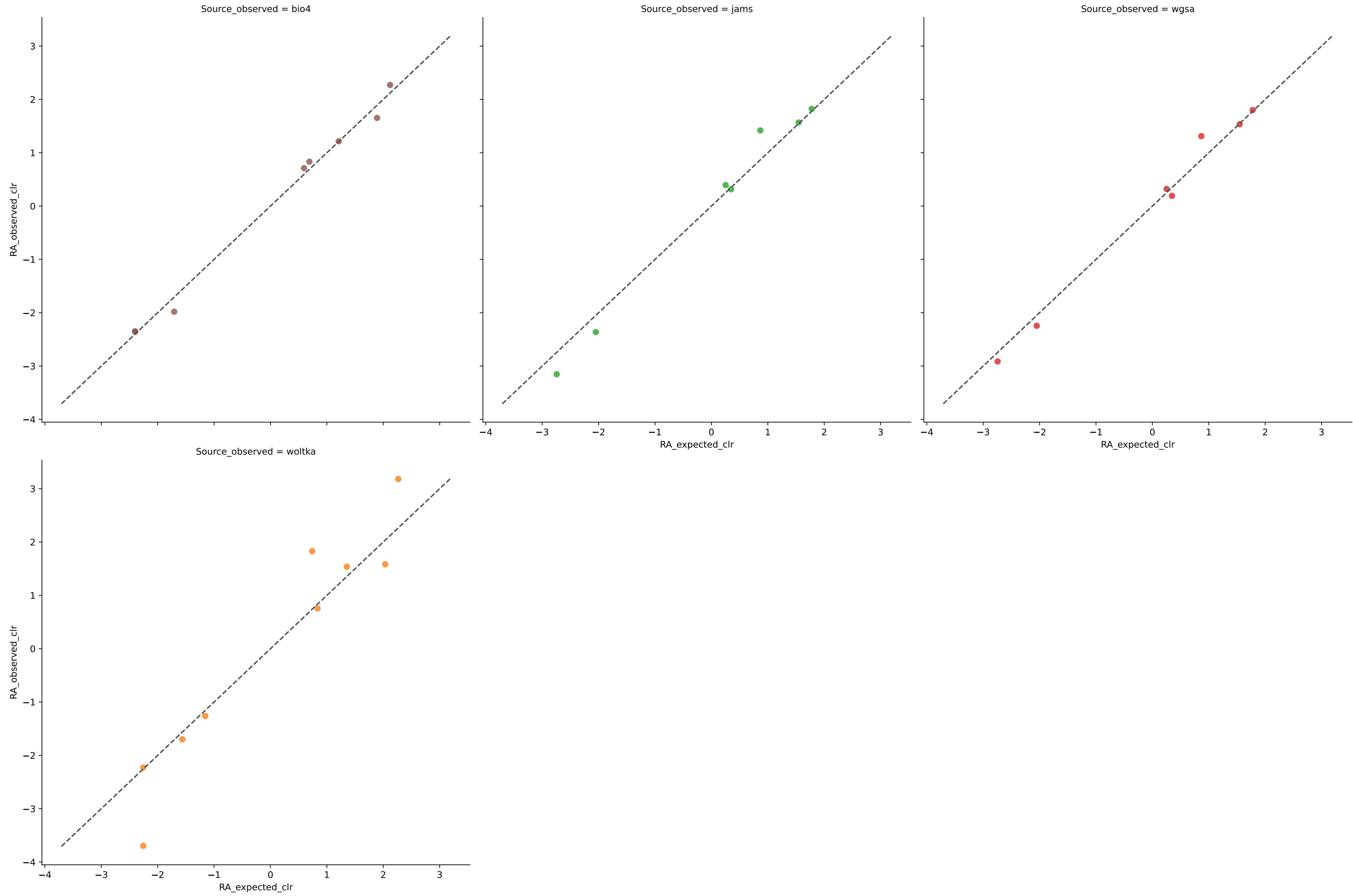


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	16	0.9954	0.0059	6.4683	0.9531	0.0130	100.0000	0.0000
jams	15	0.9996	0.0031	3.4640	0.9766	0.0047	100.0000	0.0000
wgsa	15	0.9998	0.0019	1.4092	0.9861	0.0024	100.0000	0.0000
woltka	17	0.9505	0.0143	4.1172	0.8784	0.0318	100.0000	0.0000

Bivariate Linear Regression for Sample S2 in Experiment camisimGI (Genus at filter threshold 1e-05)

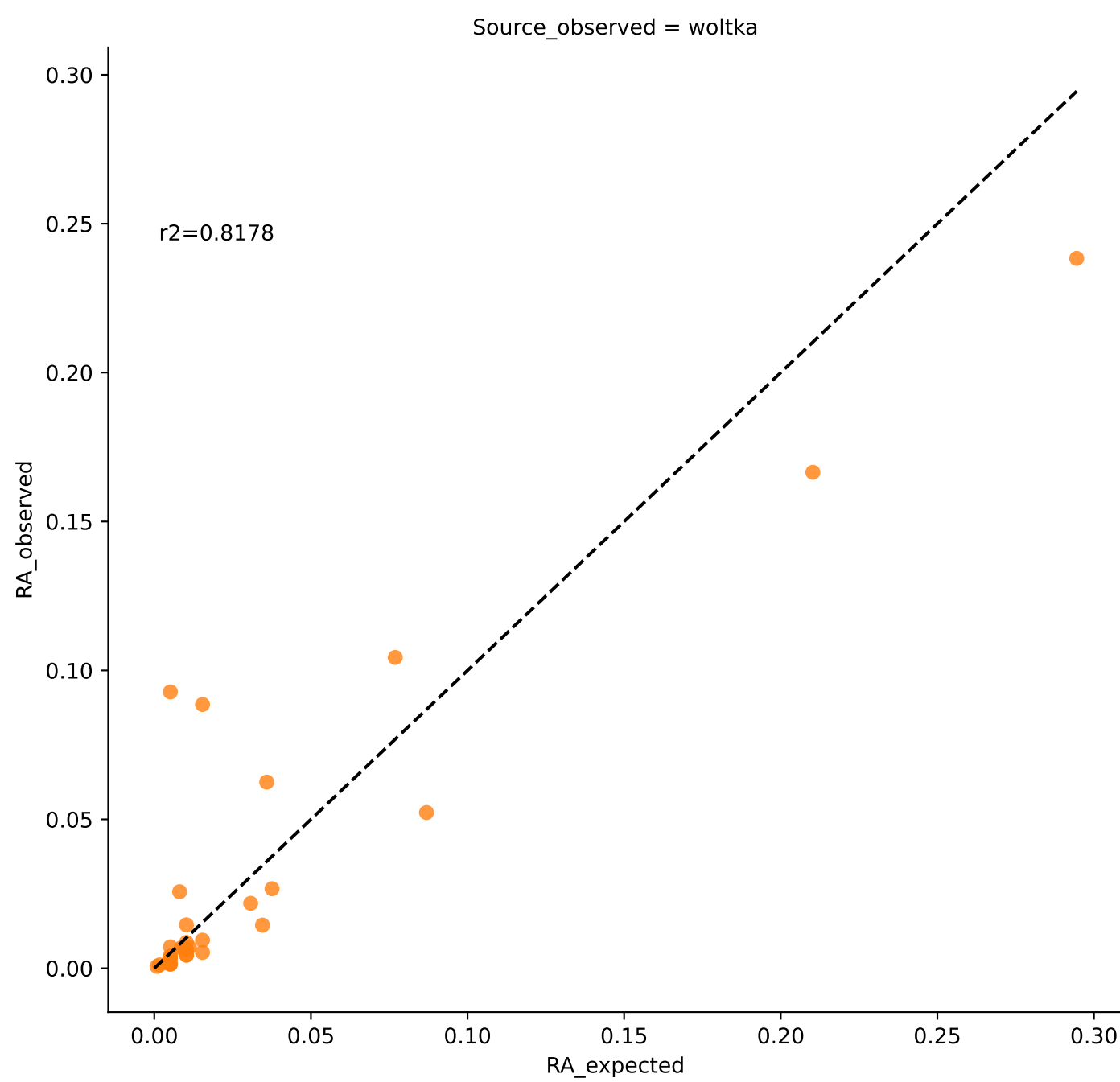
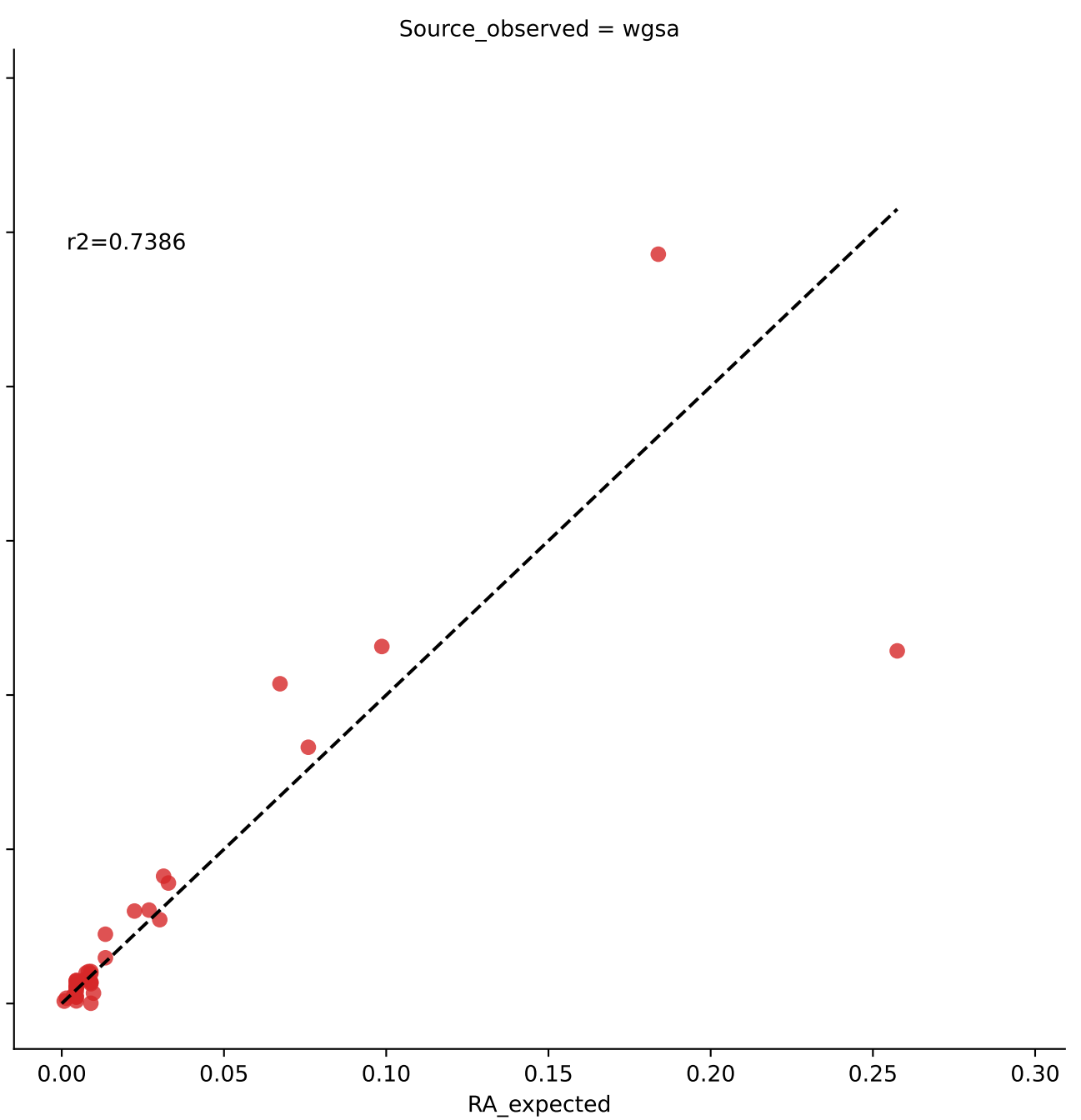
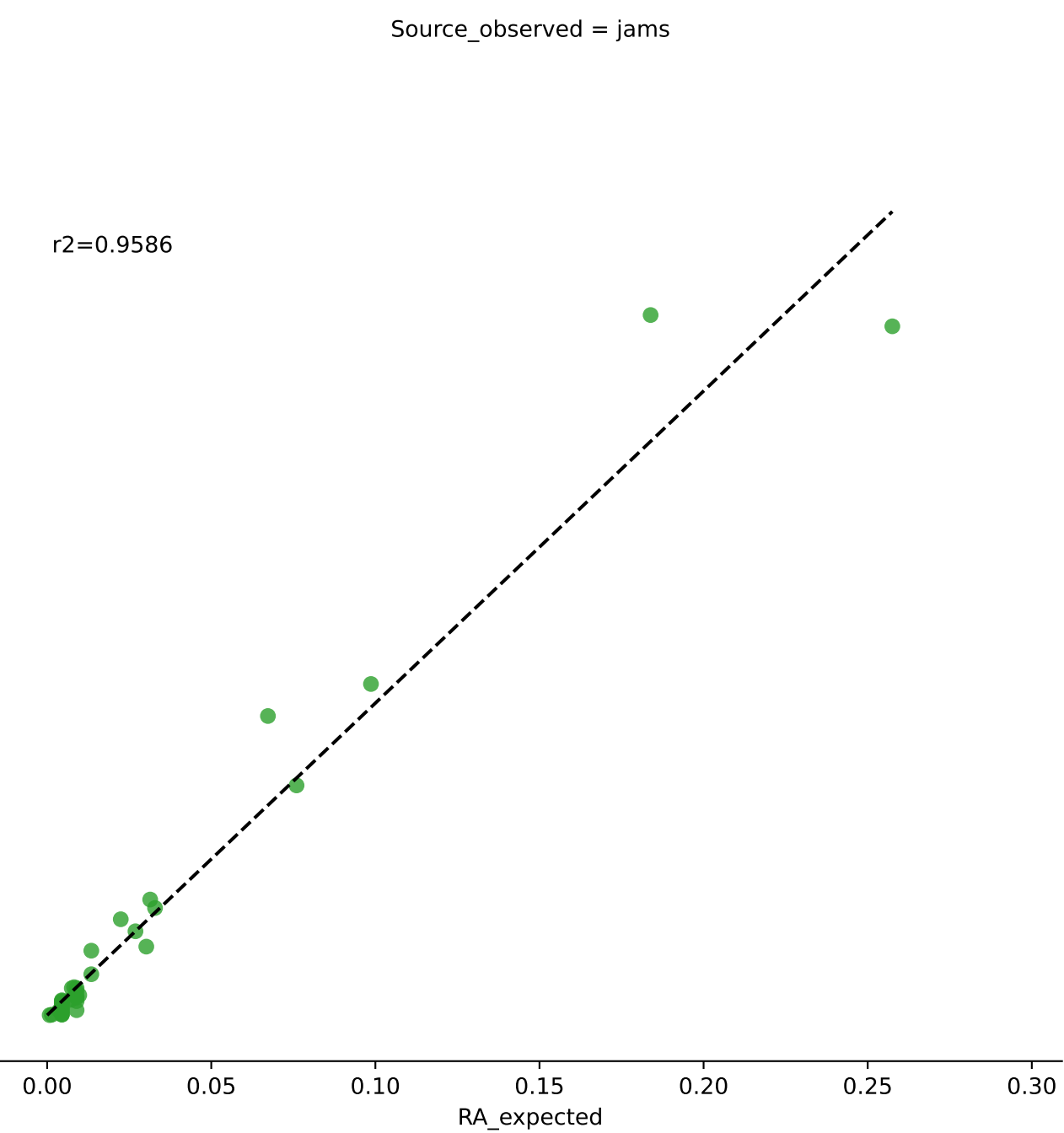
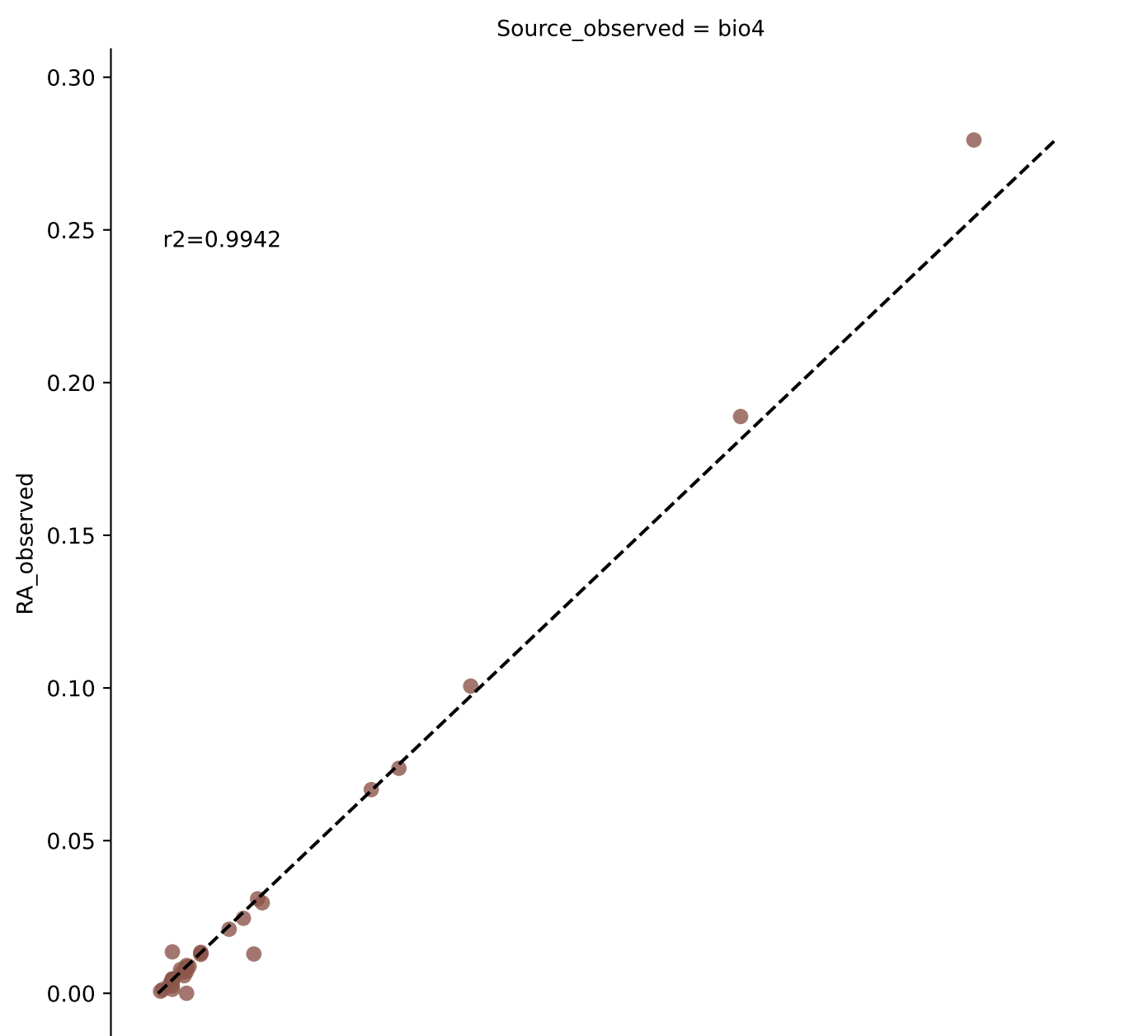


Bivariate Linear Regression for Sample S2 in Experiment camisimGI (Genus at filter threshold 1e-05)

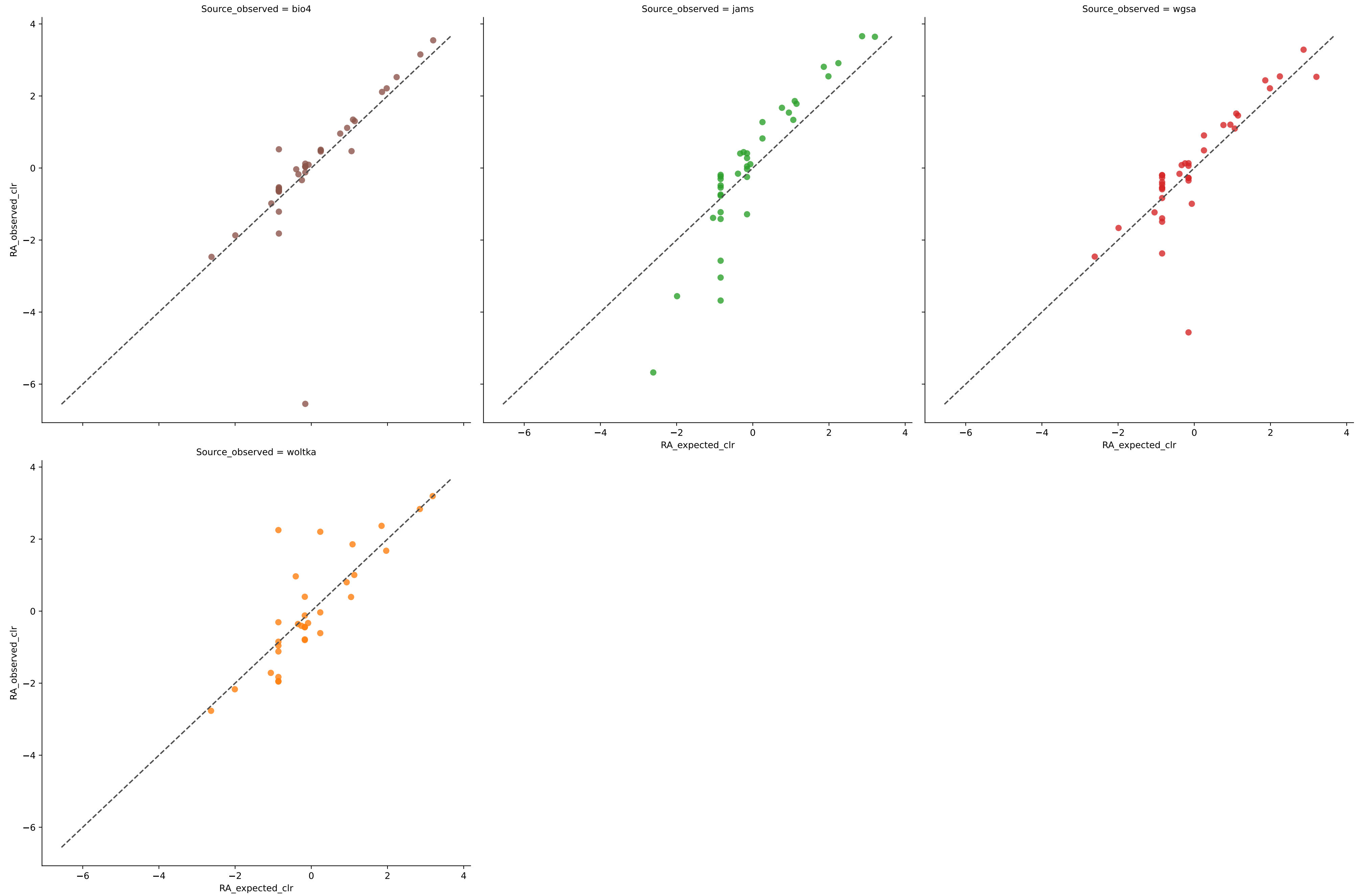


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	8	0.9494	0.0178	0.4385	0.9288	0.0302	100.0000	0.0000
jams	7	0.9317	0.0227	0.7701	0.9206	0.0343	100.0000	0.0000
wgsa	7	0.9519	0.0190	0.5411	0.9334	0.0288	100.0000	0.0000
woltka	9	0.7052	0.0592	2.0925	0.7336	0.0934	100.0000	0.0000

Bivariate Linear Regression for Sample S1 in Experiment camisimGI (Species at filter threshold 1e-05)

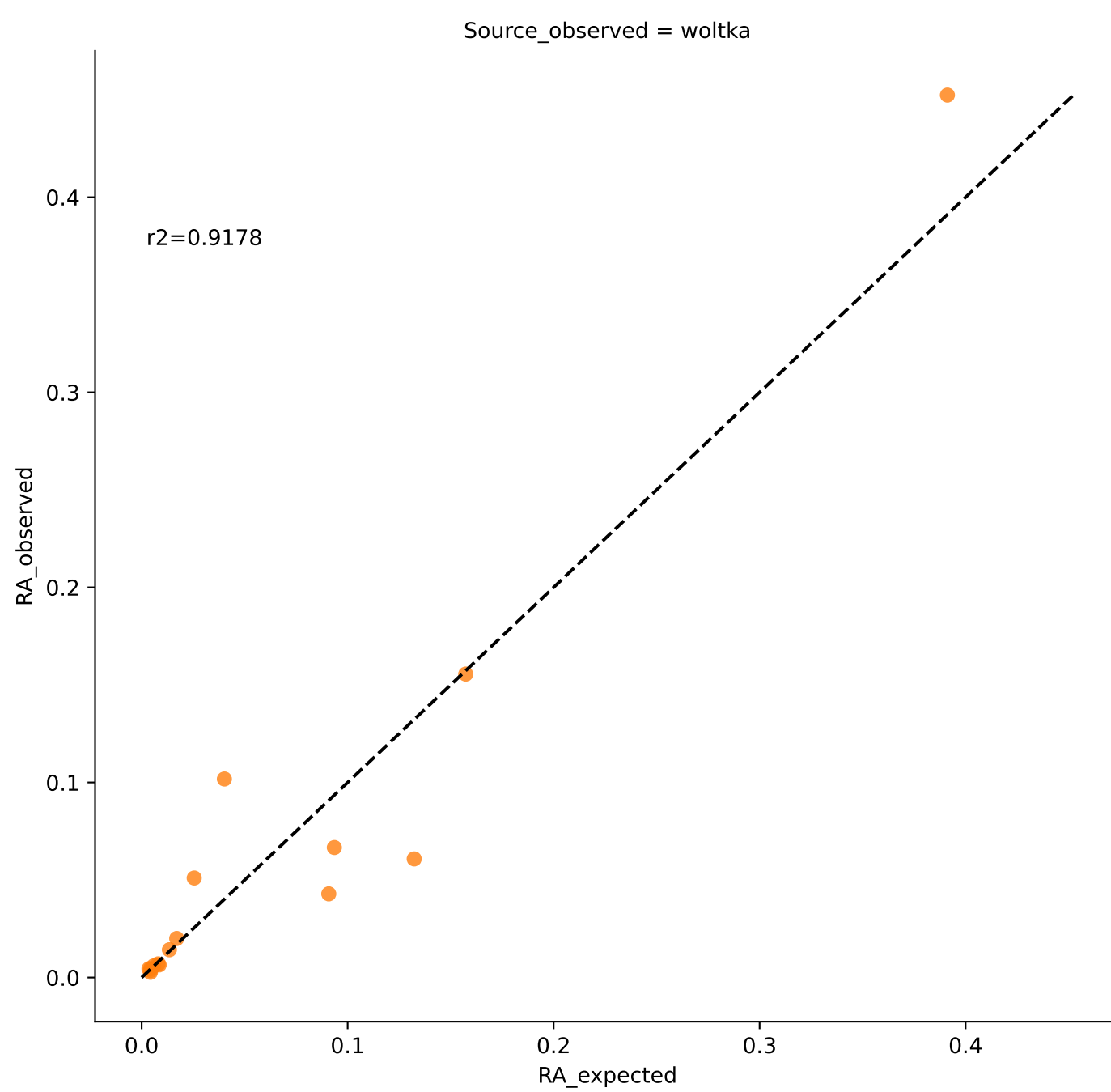
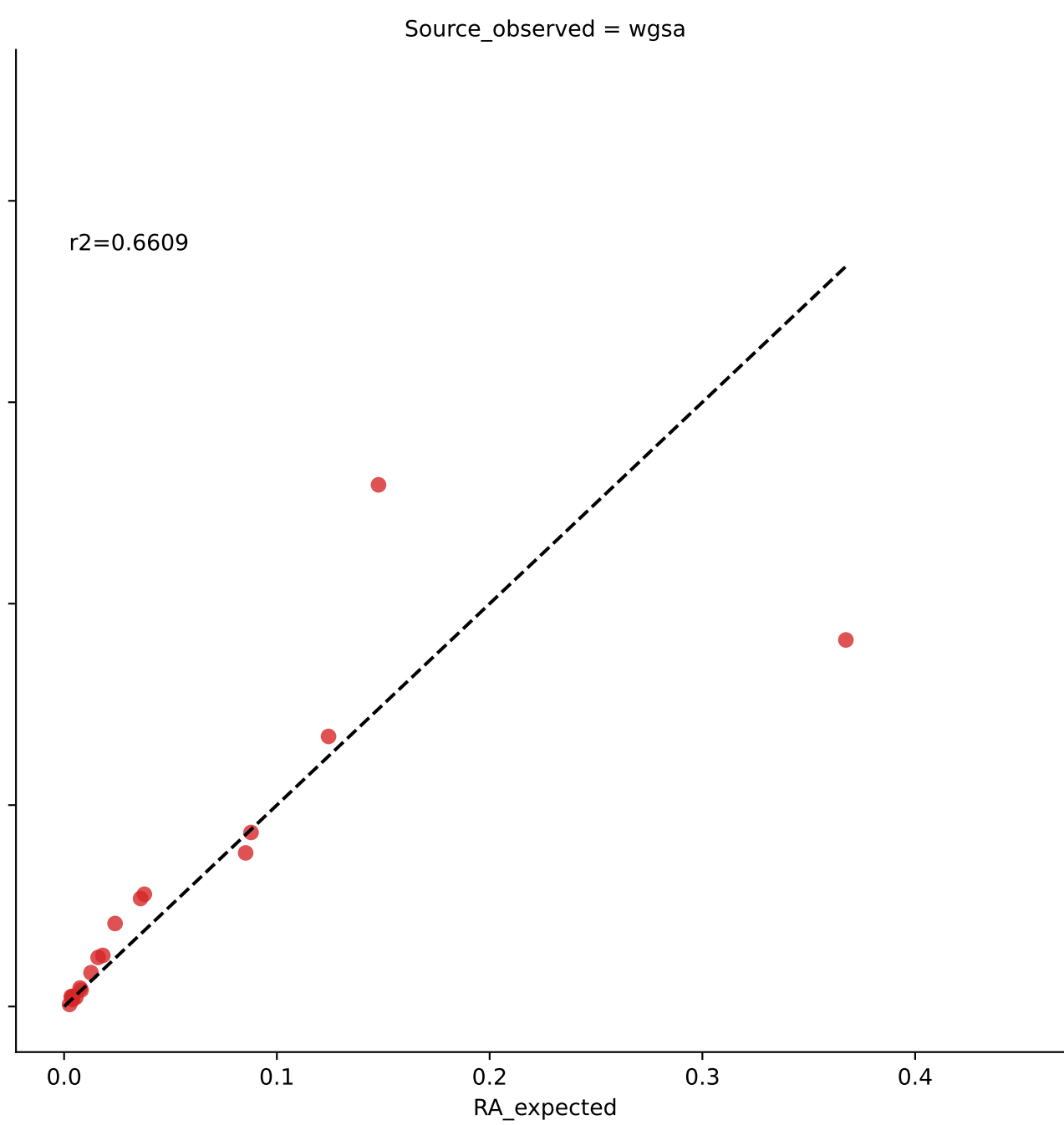
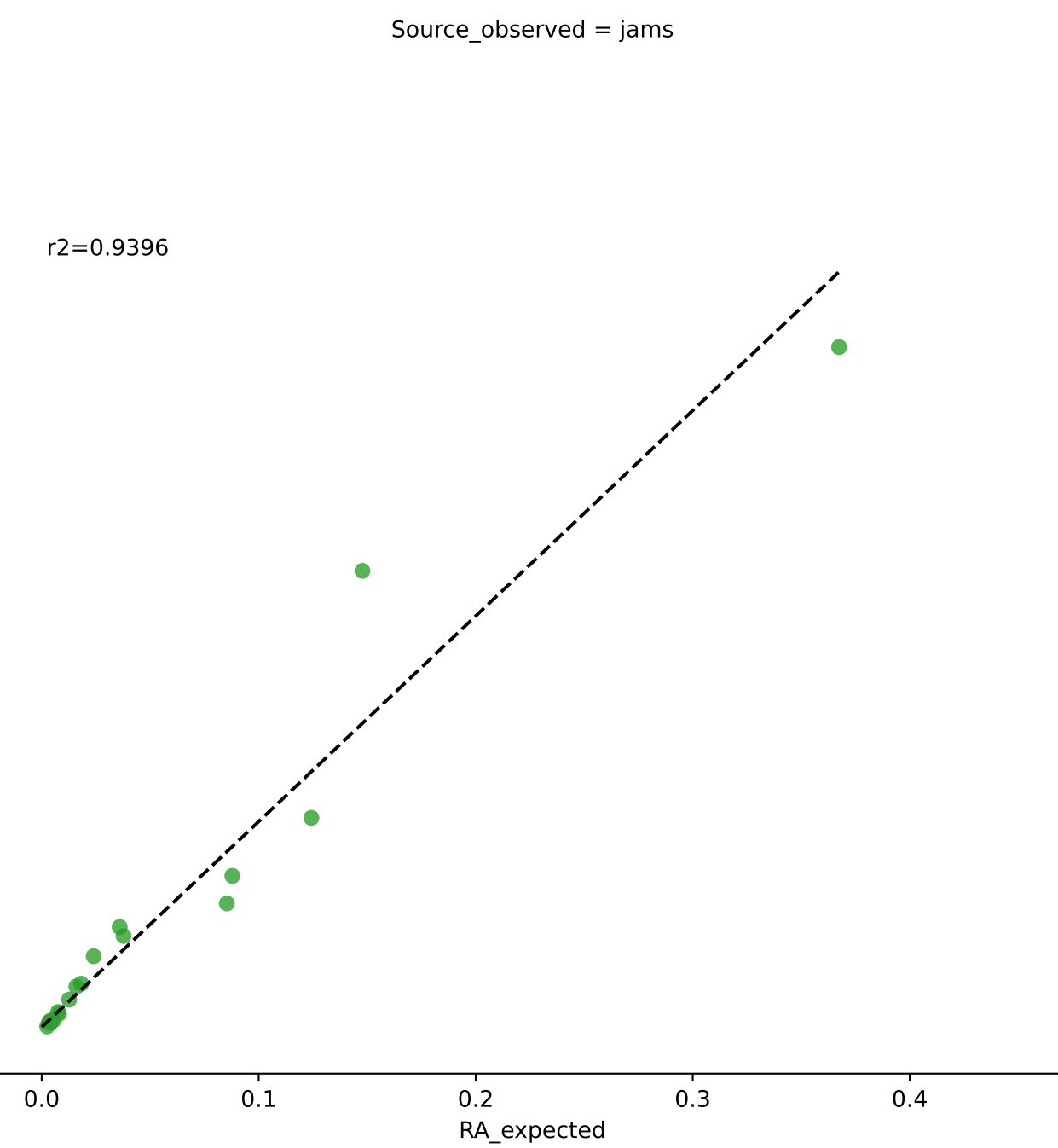
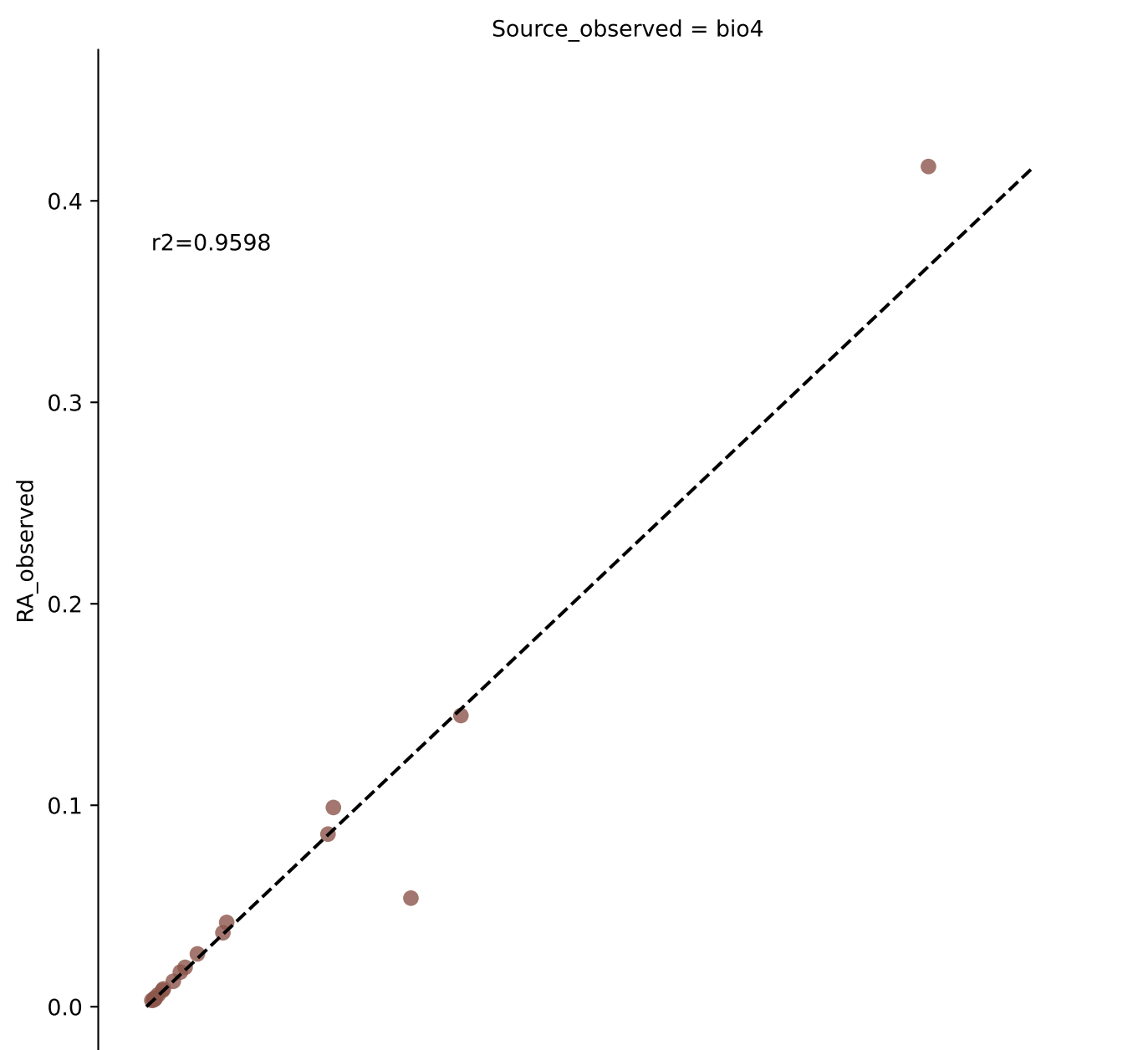


Bivariate Linear Regression for Sample S1 in Experiment camisimGI (Species at filter threshold 1e-05)

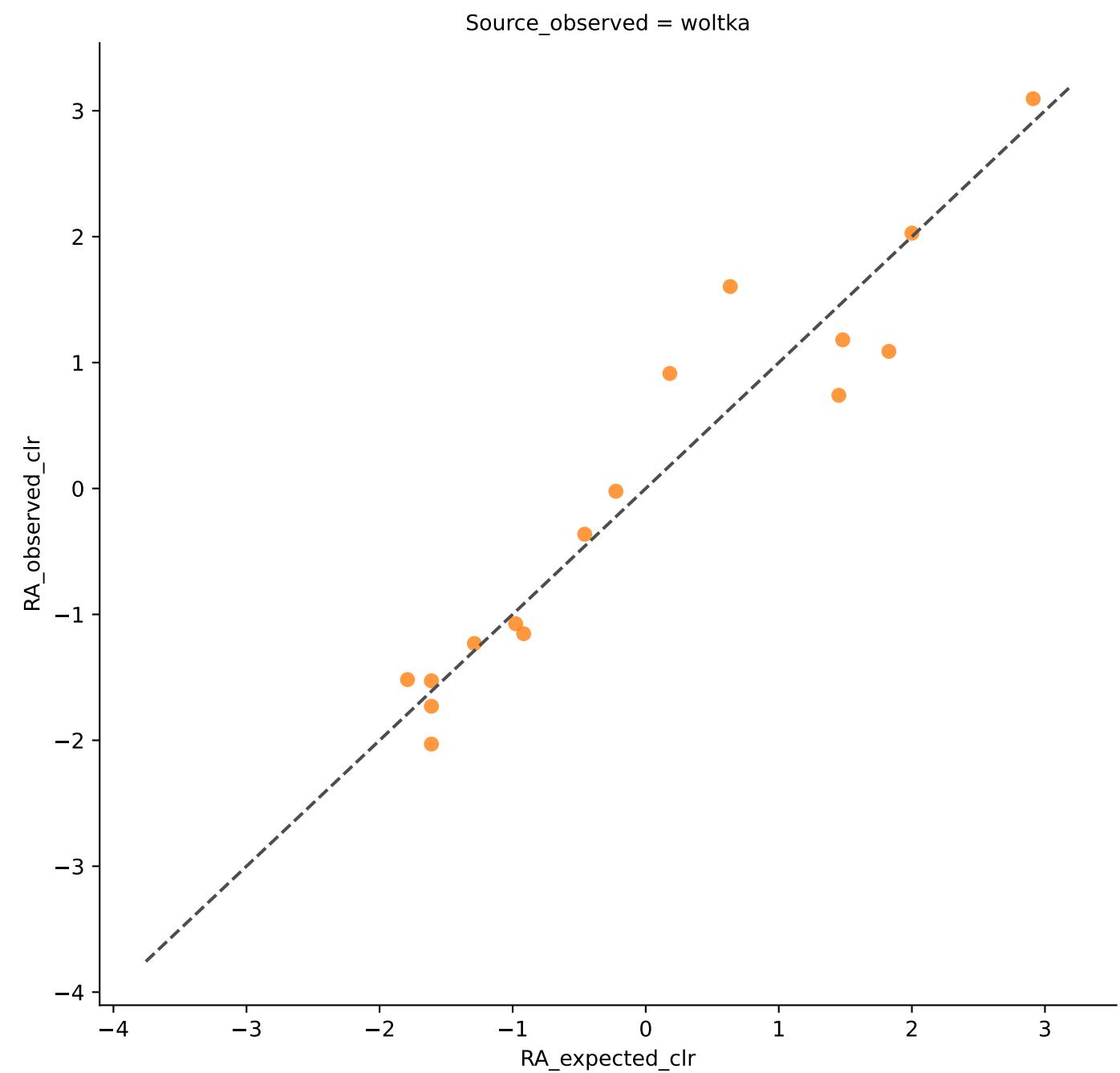
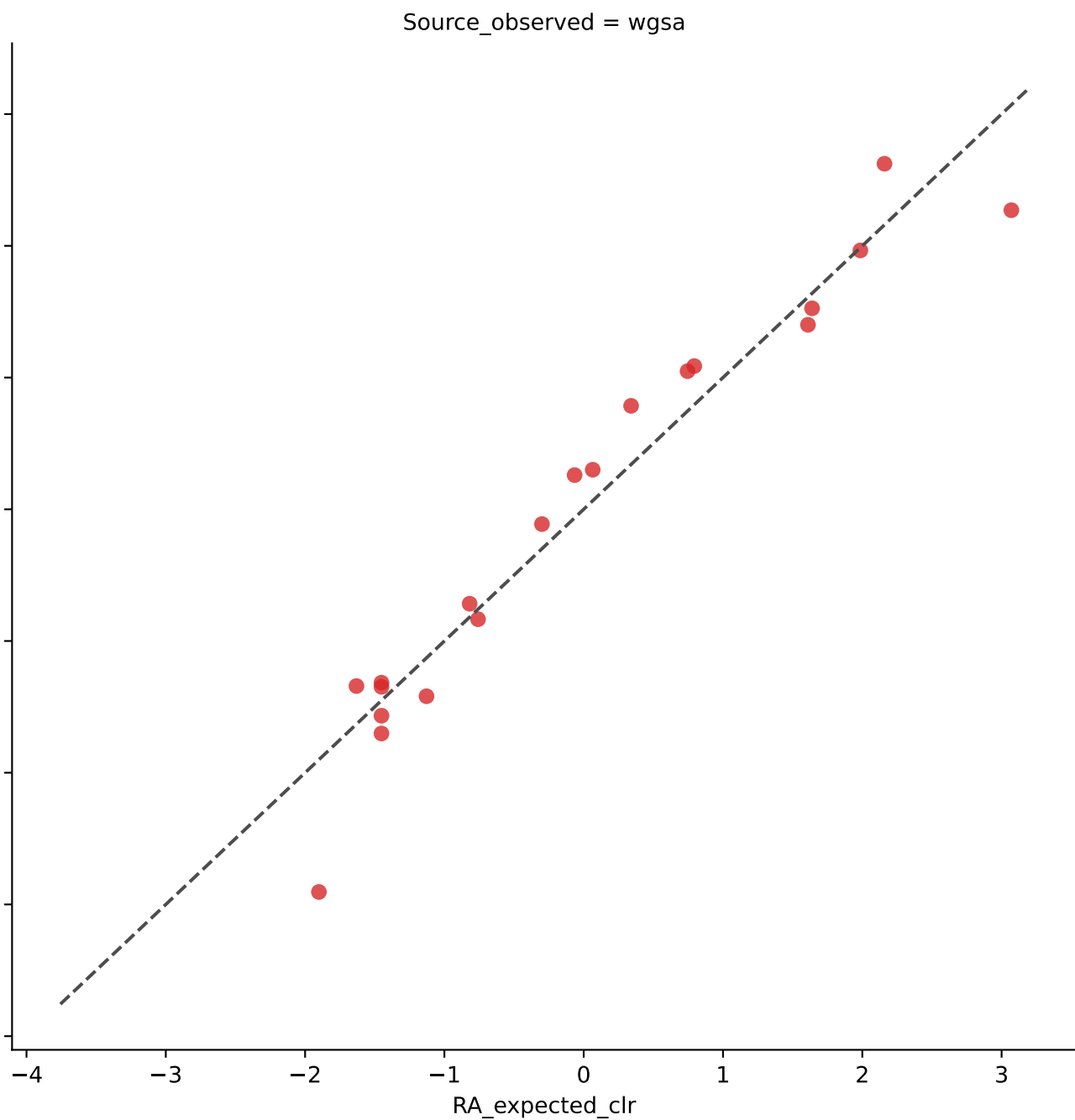
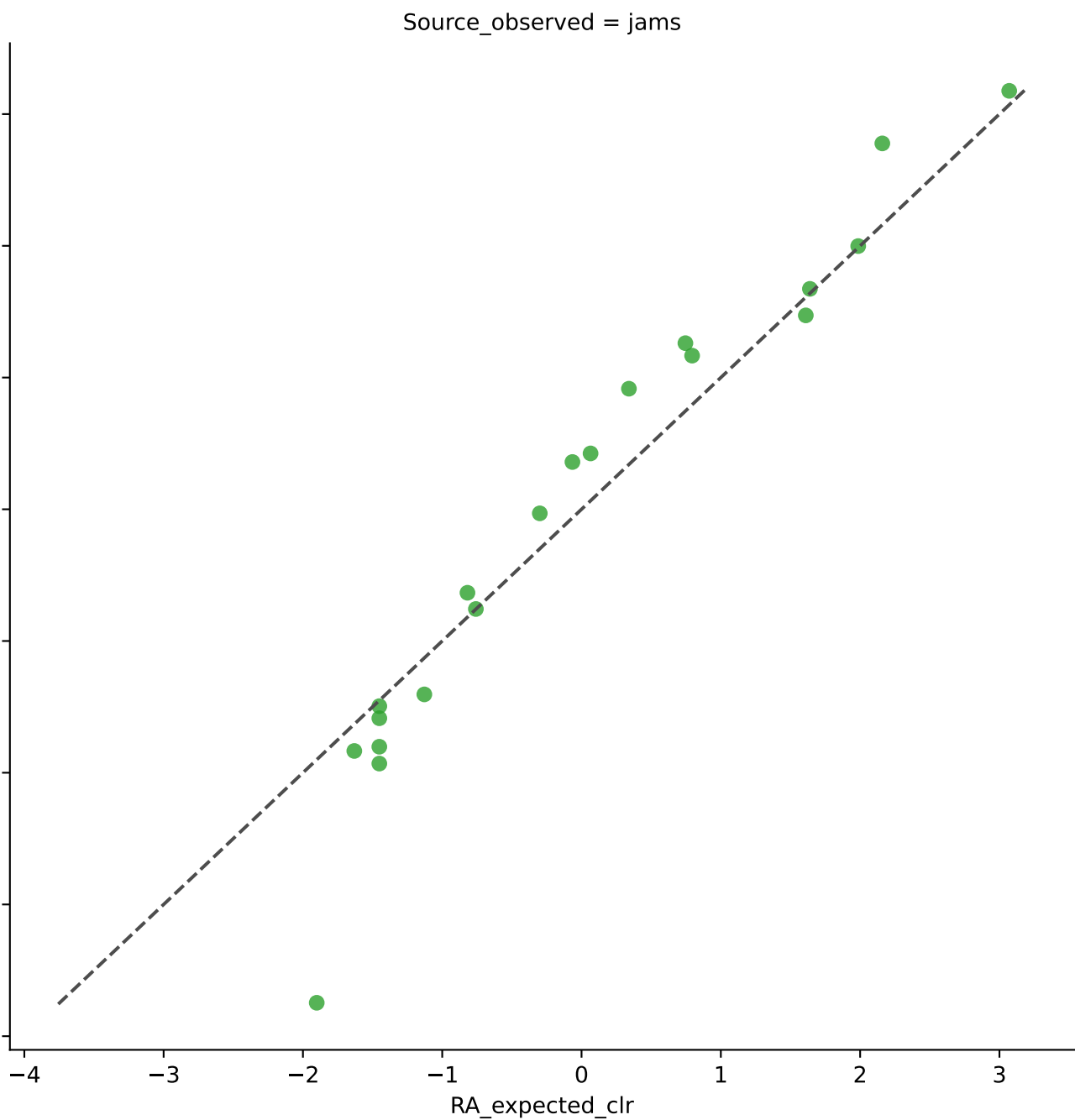
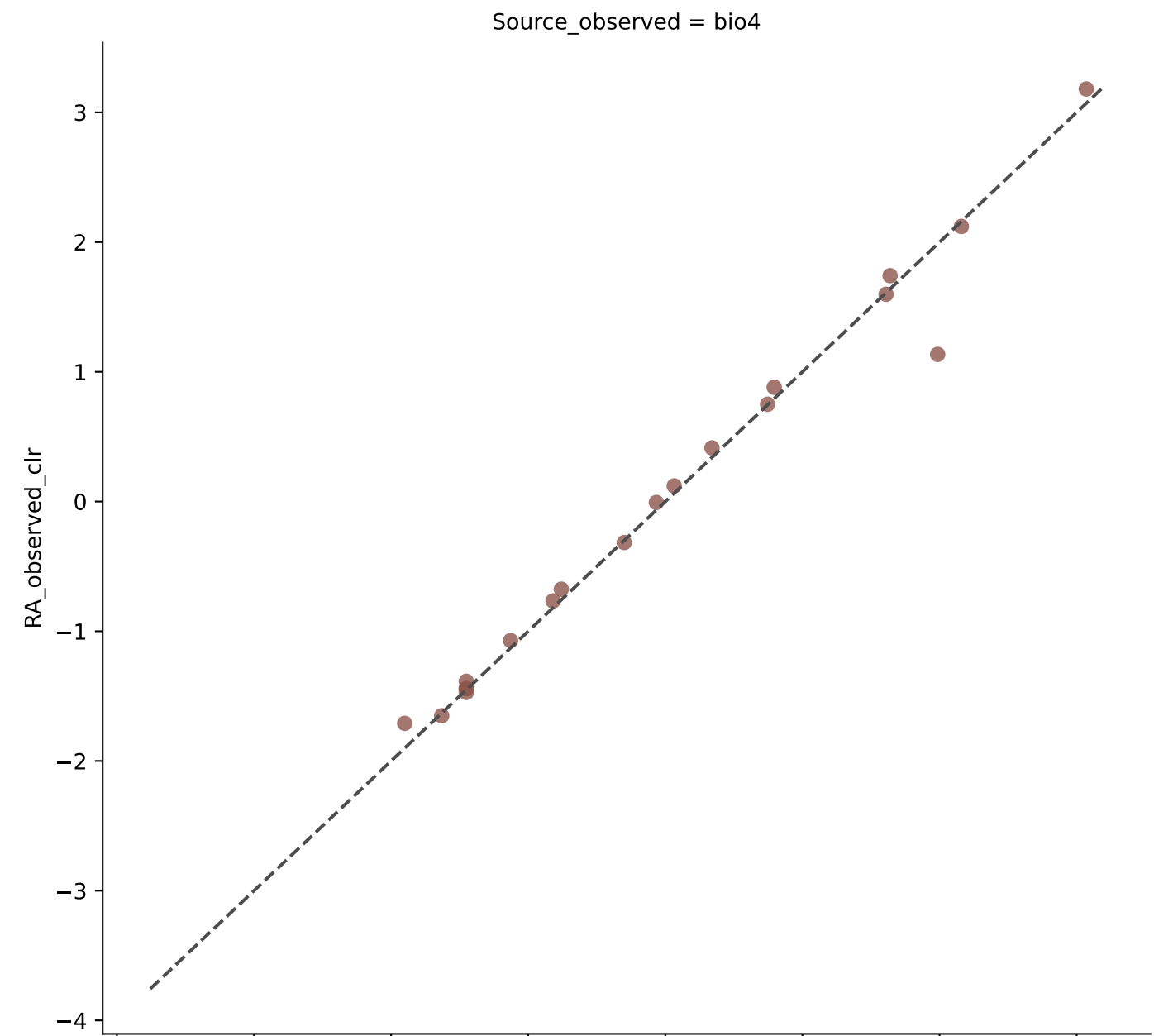


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	38	0.9942	0.0025	6.7698	0.9527	0.0056	100.0000	0.0000
jams	37	0.9586	0.0055	6.2135	0.8985	0.0108	100.0000	0.0000
wgsa	37	0.7386	0.0096	5.2876	0.8219	0.0266	100.0000	0.0000
woltka	32	0.8178	0.0149	4.8097	0.7611	0.0262	100.0000	0.0000

Bivariate Linear Regression for Sample S2 in Experiment camsimGI (Species at filter threshold 1e-05)

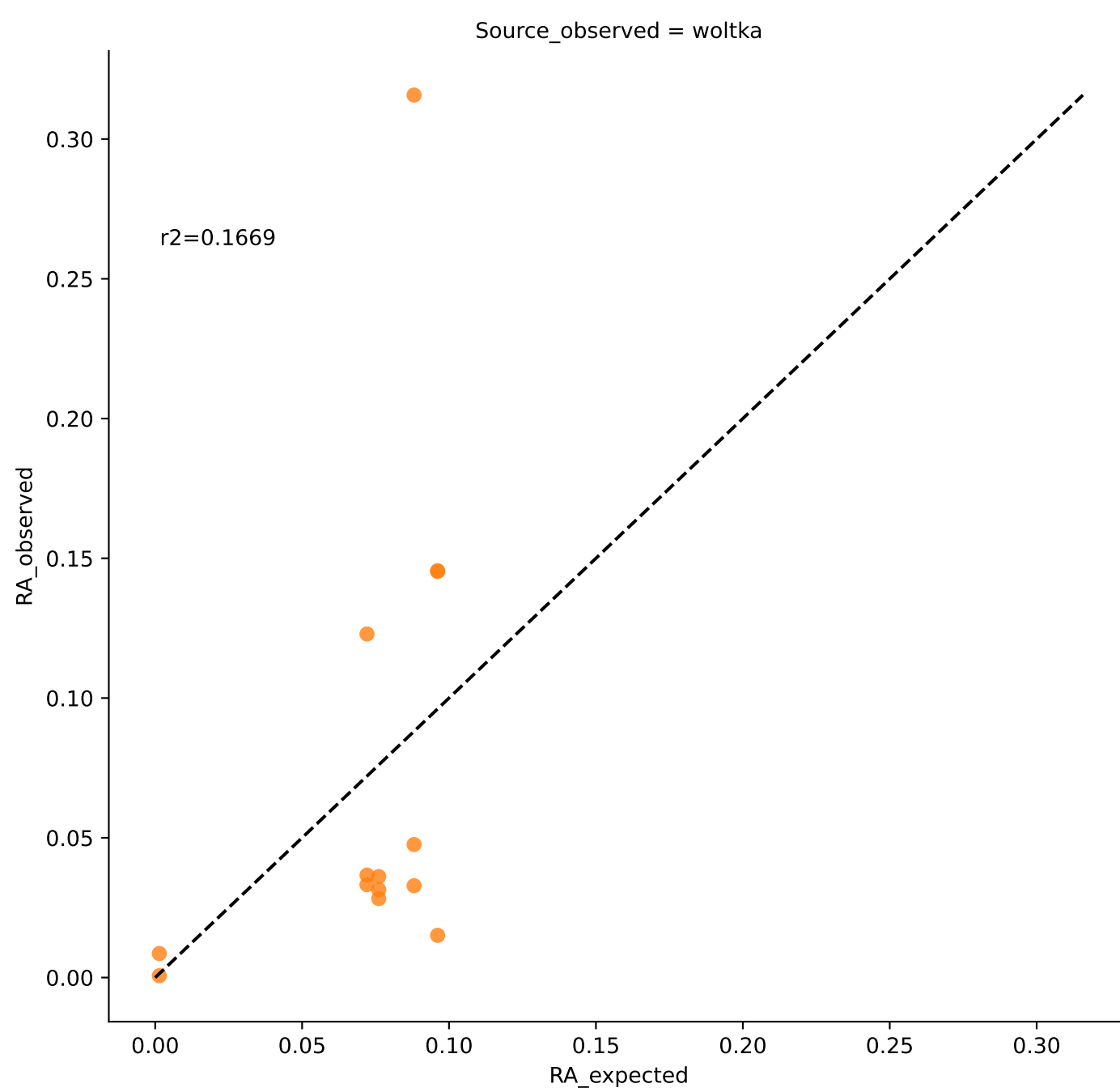
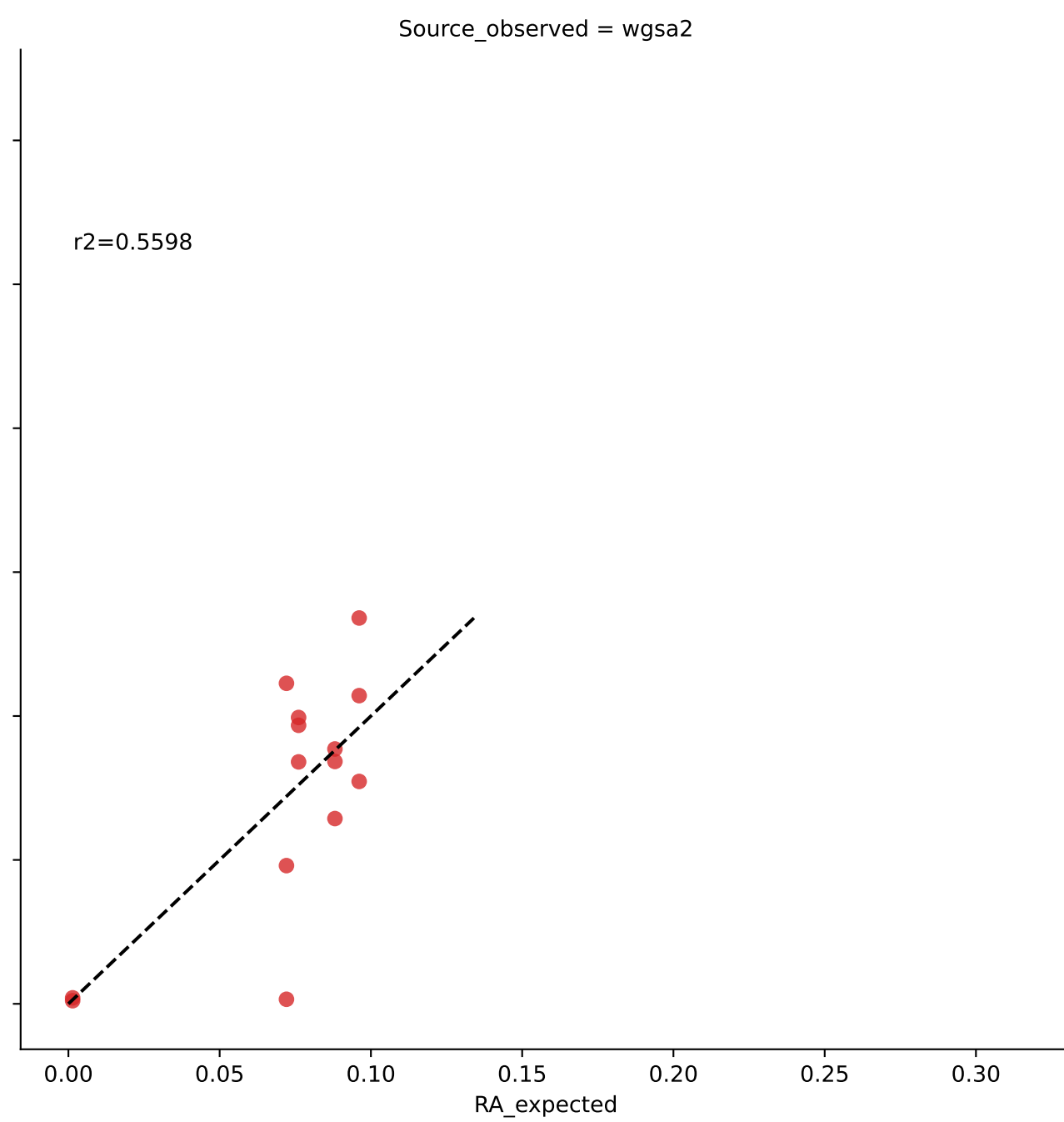
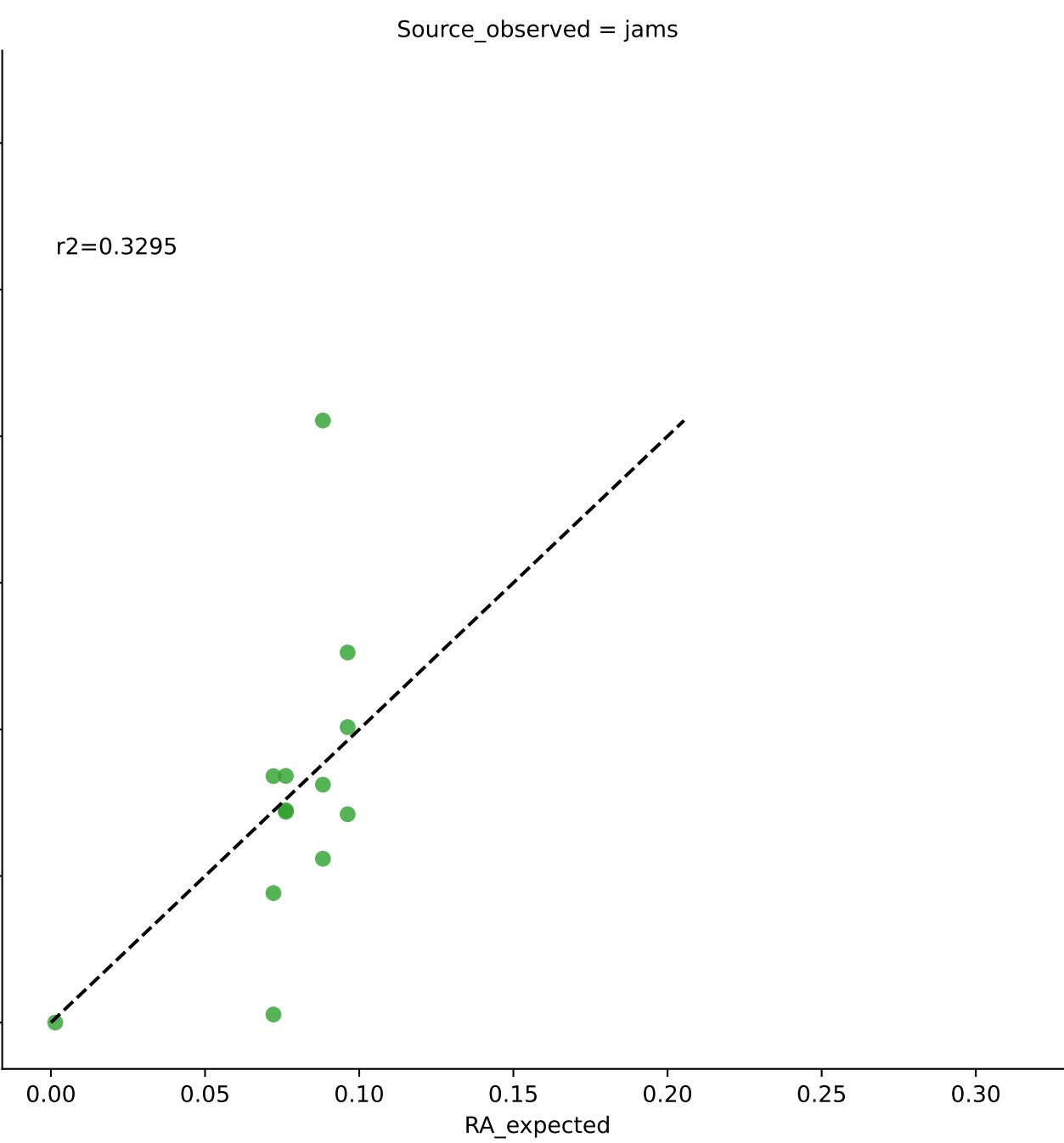
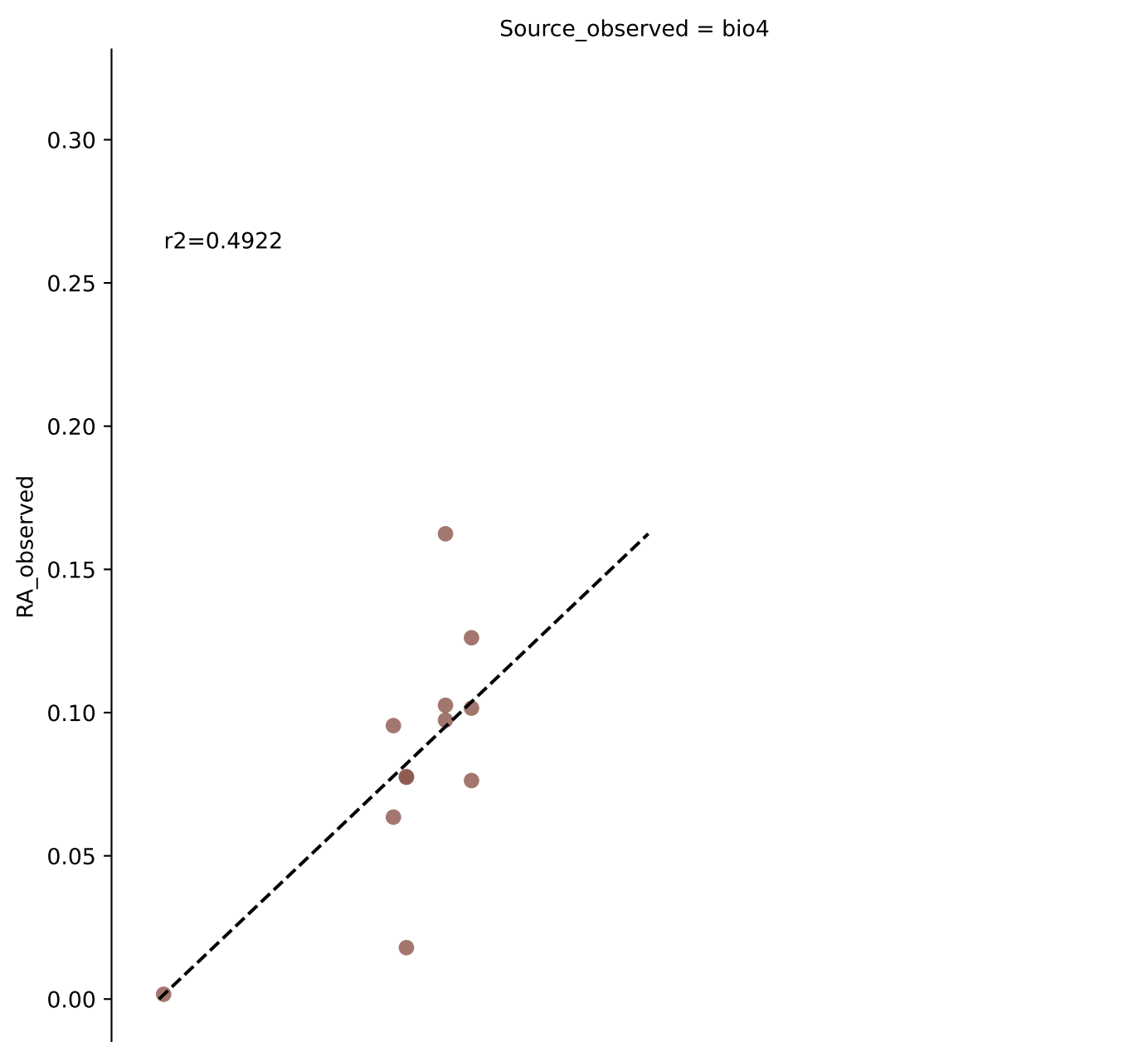


Bivariate Linear Regression for Sample S2 in Experiment camisimGI (Species at filter threshold 1e-05)

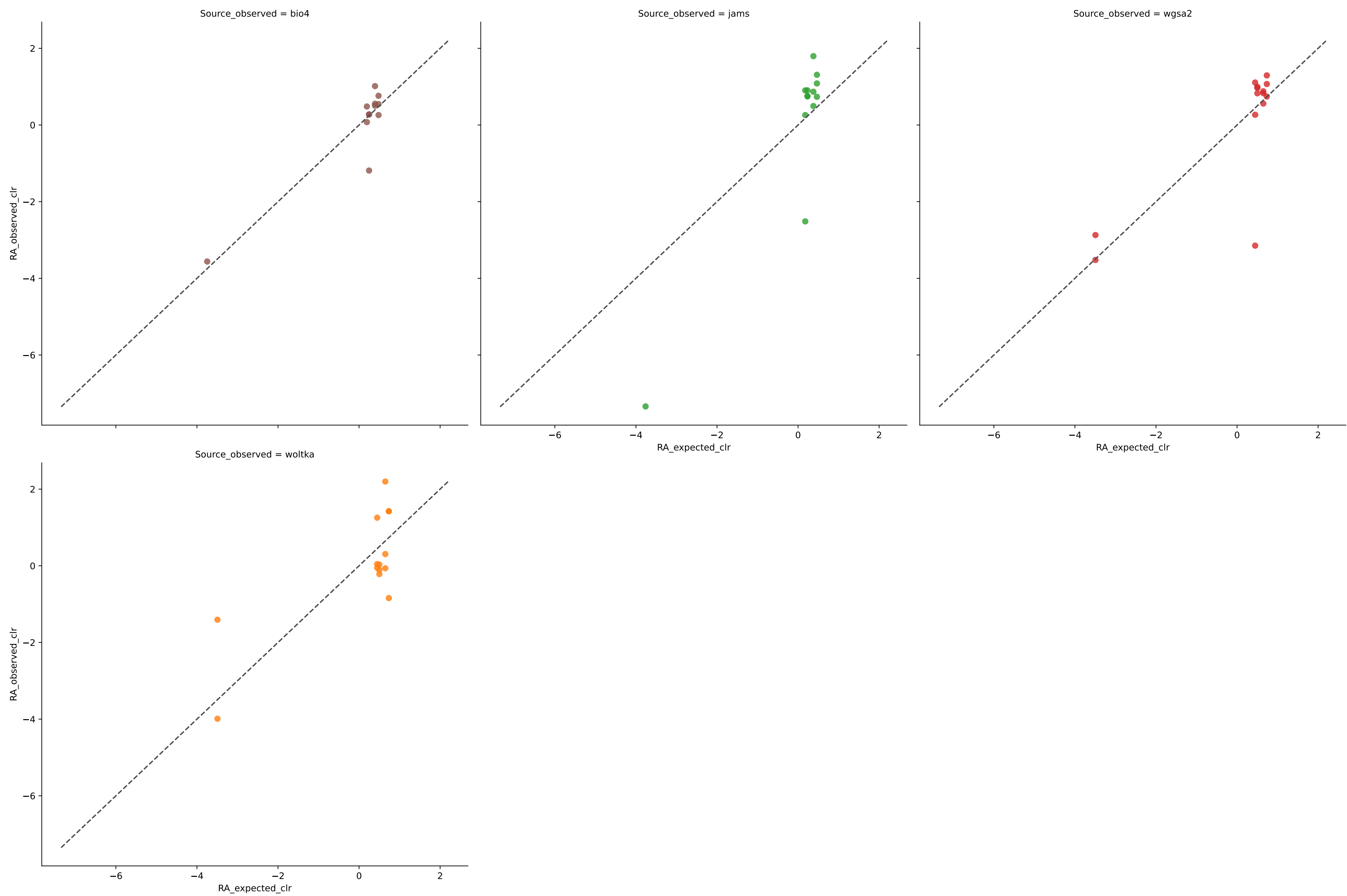


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	20	0.9598	0.0074	0.9077	0.9264	0.0195	100.0000	0.0000
jams	20	0.9396	0.0112	2.3368	0.8885	0.0207	100.0000	0.0000
wgsa	20	0.6609	0.0199	1.6681	0.8009	0.0490	100.0000	0.0000
woltka	16	0.9178	0.0192	1.7436	0.8467	0.0320	100.0000	0.0000

Bivariate Linear Regression for Sample EG in Experiment nist (Genus at filter threshold 1e-05)

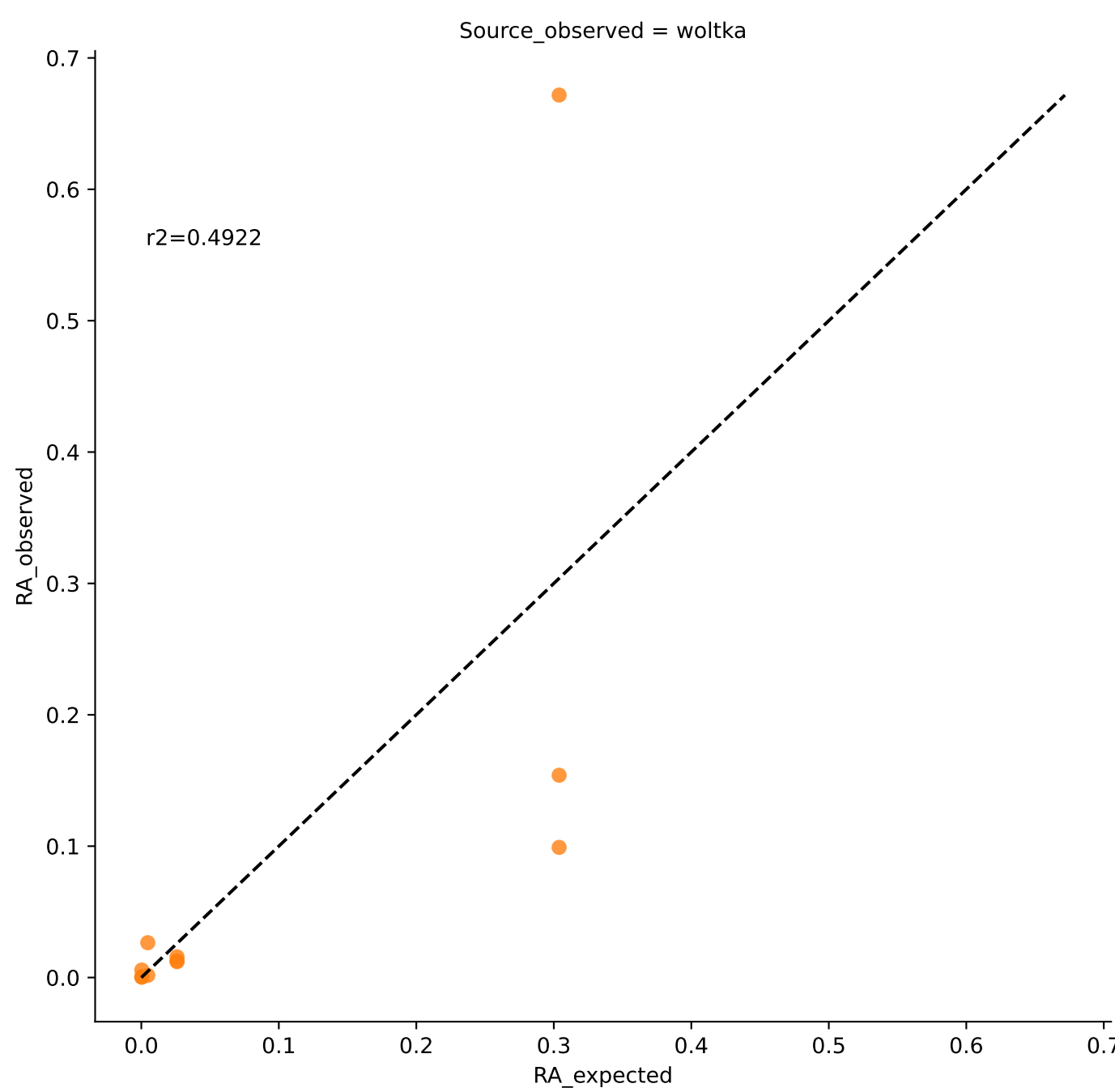
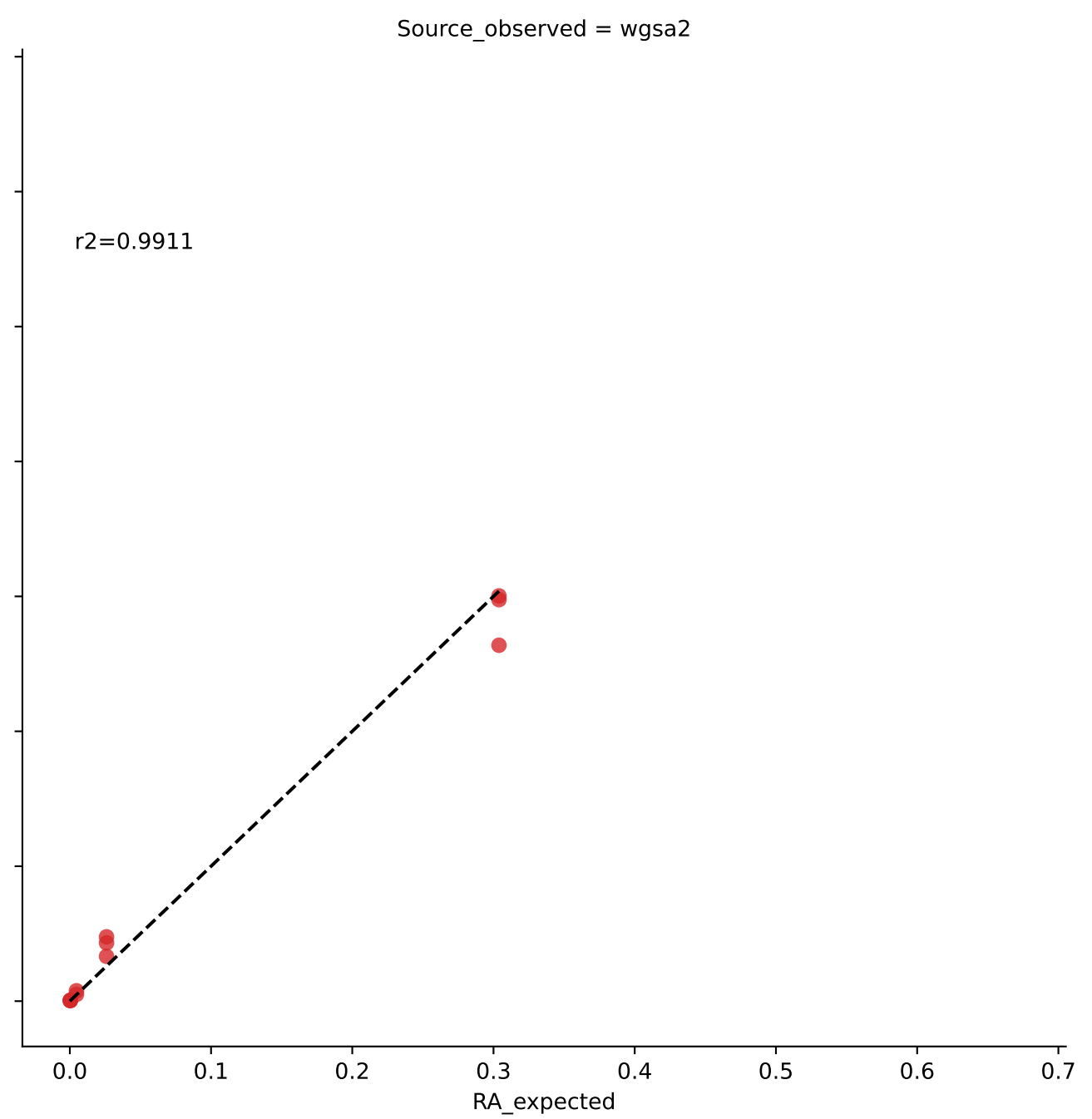
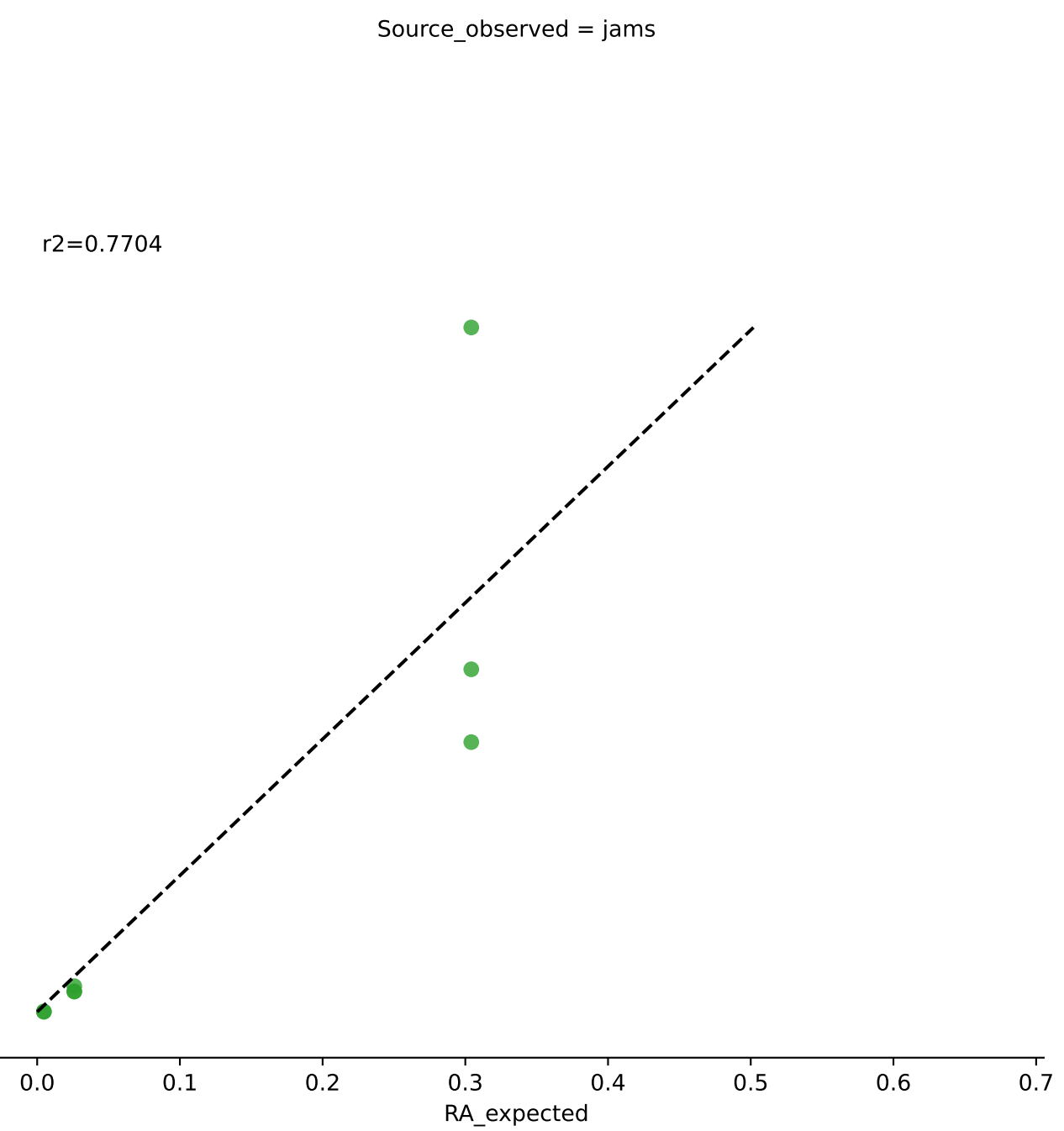
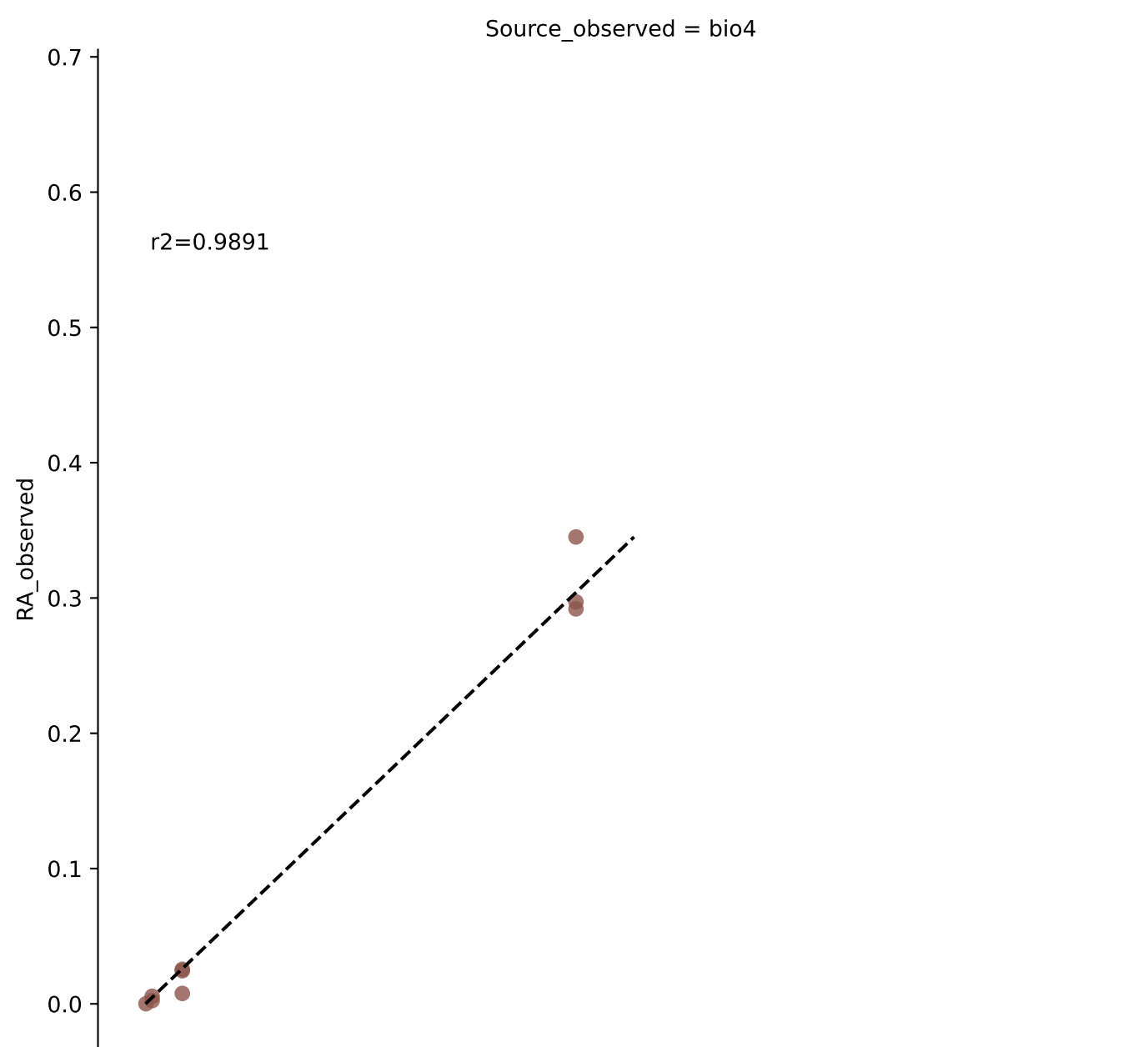


Bivariate Linear Regression for Sample EG in Experiment nist (Genus at filter threshold 1e-05)

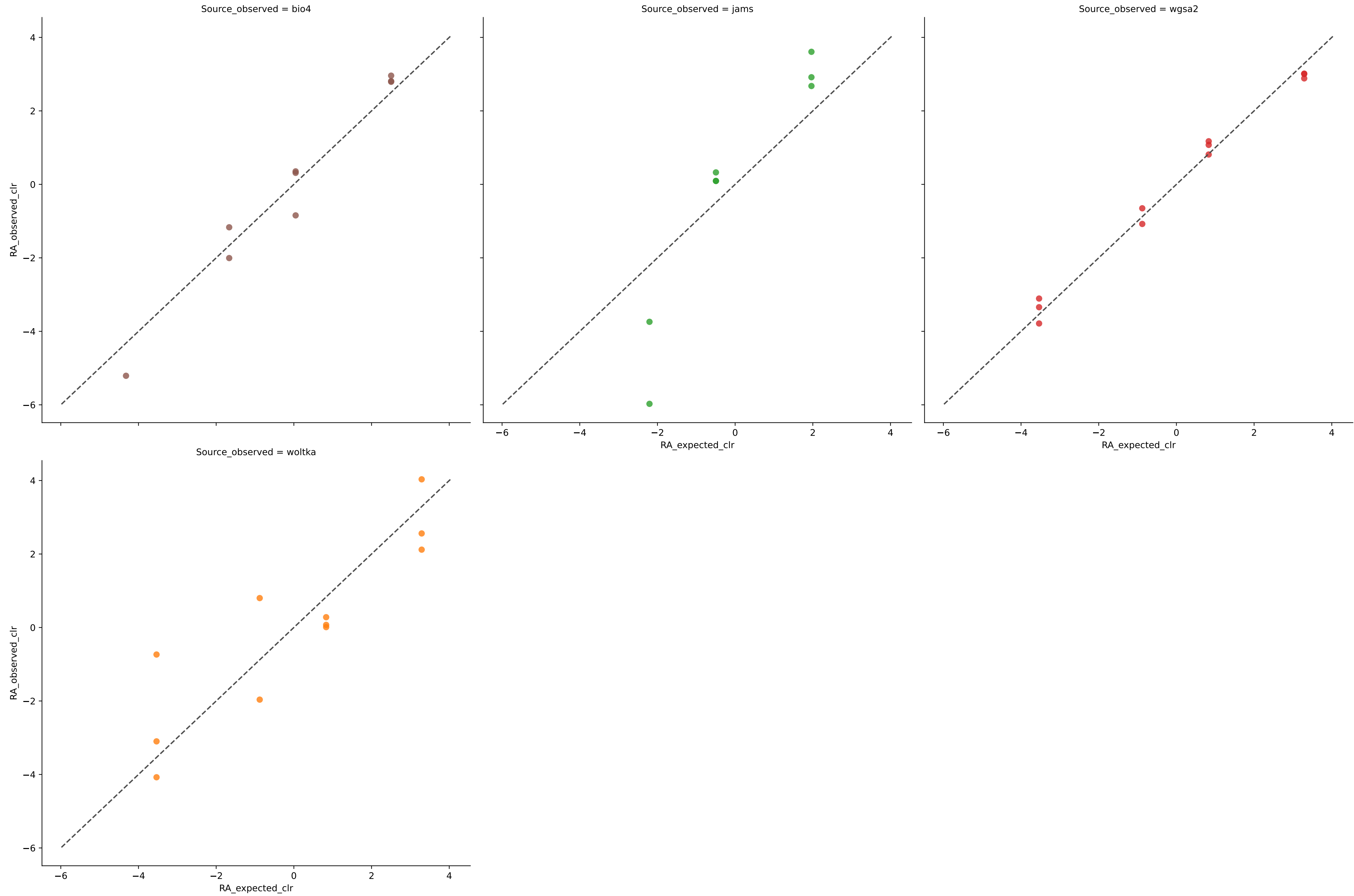


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	12	0.4922	0.0195	1.6584	0.8827	0.0296	100.0000	0.0000
jams	13	0.3295	0.0264	5.0005	0.8285	0.0413	100.0000	0.0000
wgsa2	14	0.5598	0.0202	3.8550	0.8586	0.0276	100.0000	0.0000
woltka	14	0.1669	0.0549	3.6377	0.6157	0.0752	100.0000	0.0000

Bivariate Linear Regression for Sample MIX-A in Experiment nist (Genus at filter threshold 1e-05)

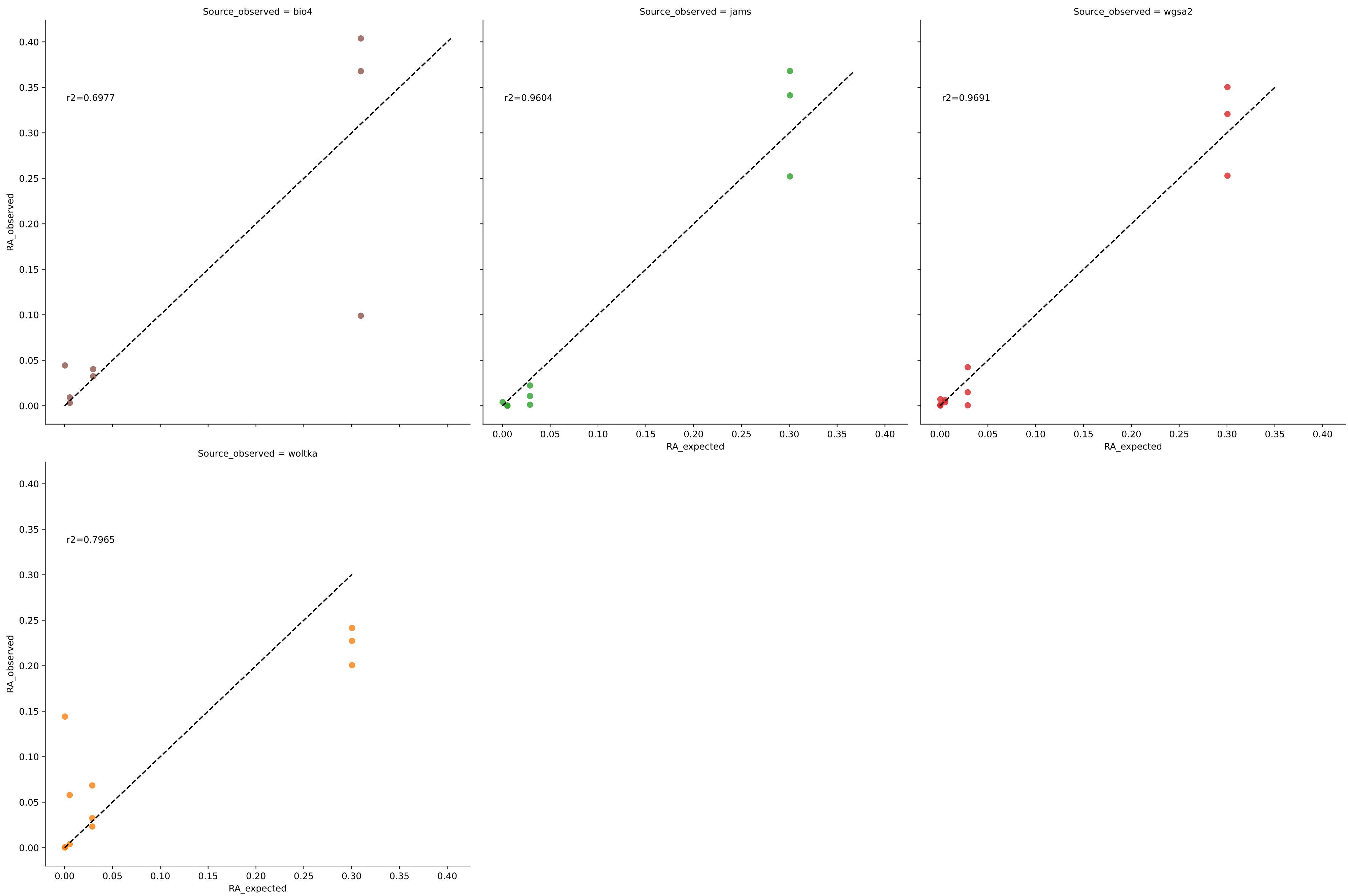


Bivariate Linear Regression for Sample MIX-A in Experiment nist (Genus at filter threshold 1e-05)

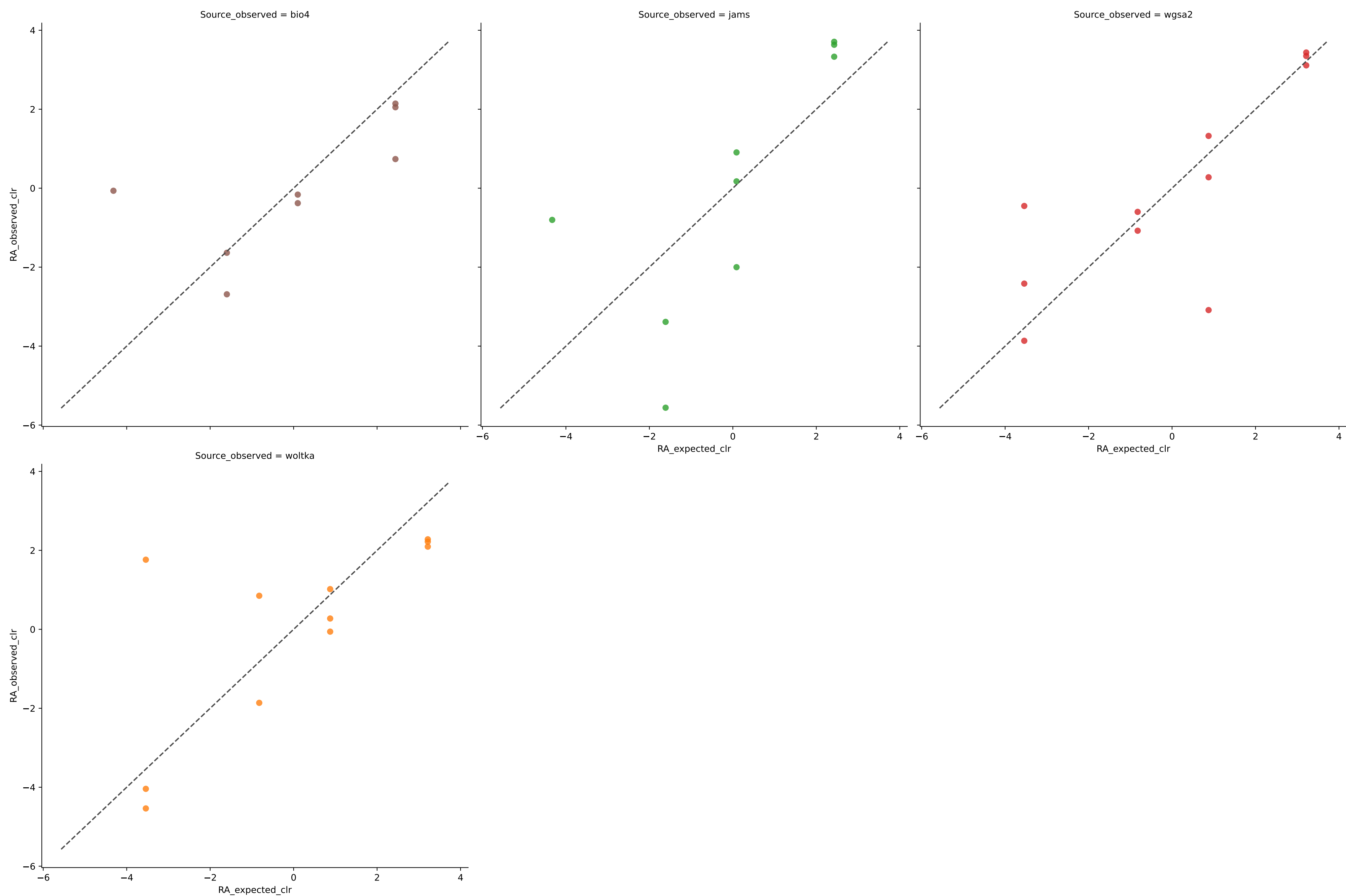


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	9	0.9891	0.0093	1.5789	0.9581	0.0157	100.0000	0.0000
jams	8	0.7704	0.0494	4.6943	0.8023	0.0818	100.0000	0.0000
wgsa2	11	0.9911	0.0091	0.9344	0.9501	0.0150	100.0000	0.0000
woltka	11	0.4922	0.0719	4.0410	0.6048	0.1351	100.0000	0.0000

Bivariate Linear Regression for Sample MIX-B in Experiment nist (Genus at filter threshold 1e-05)

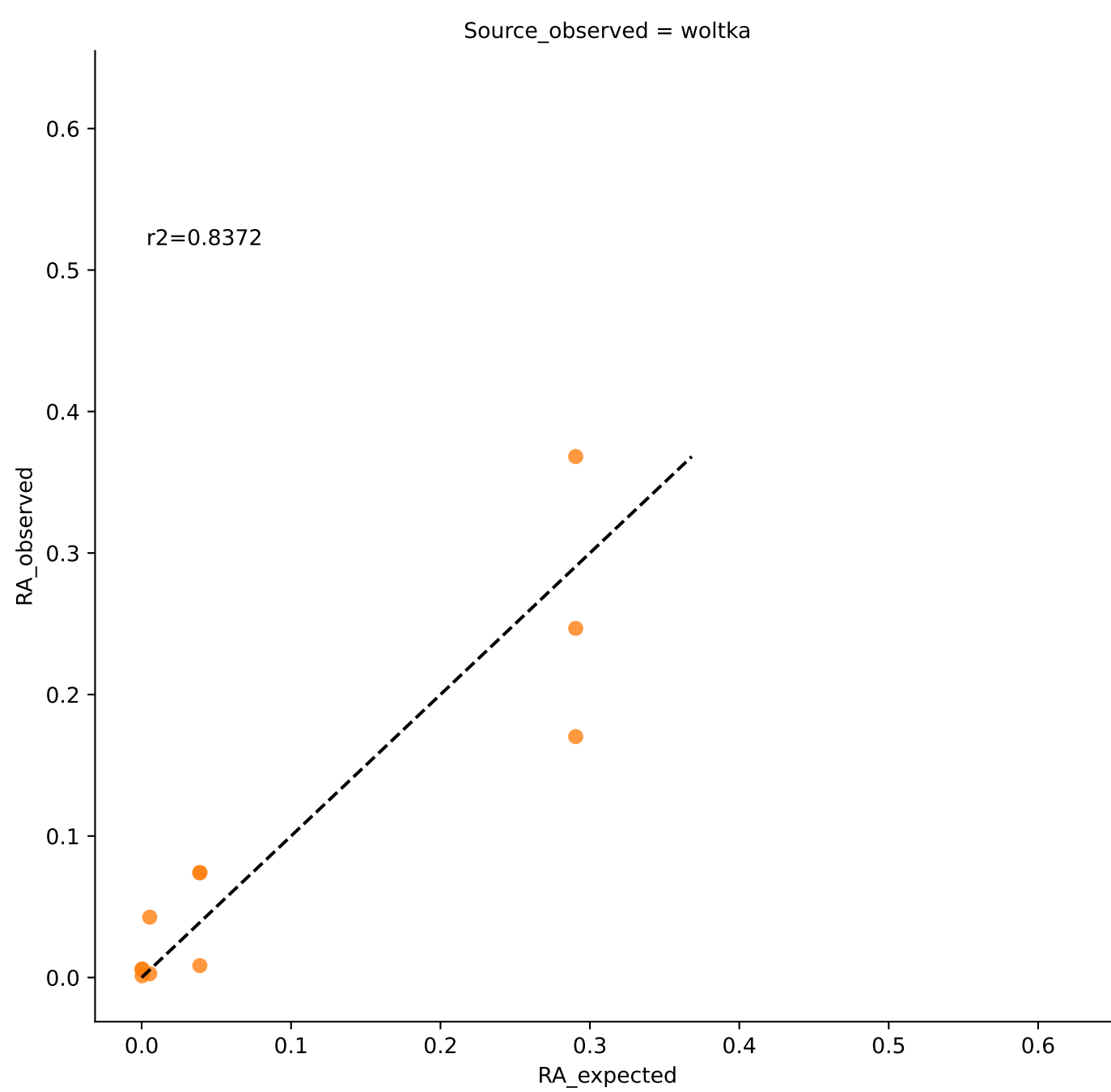
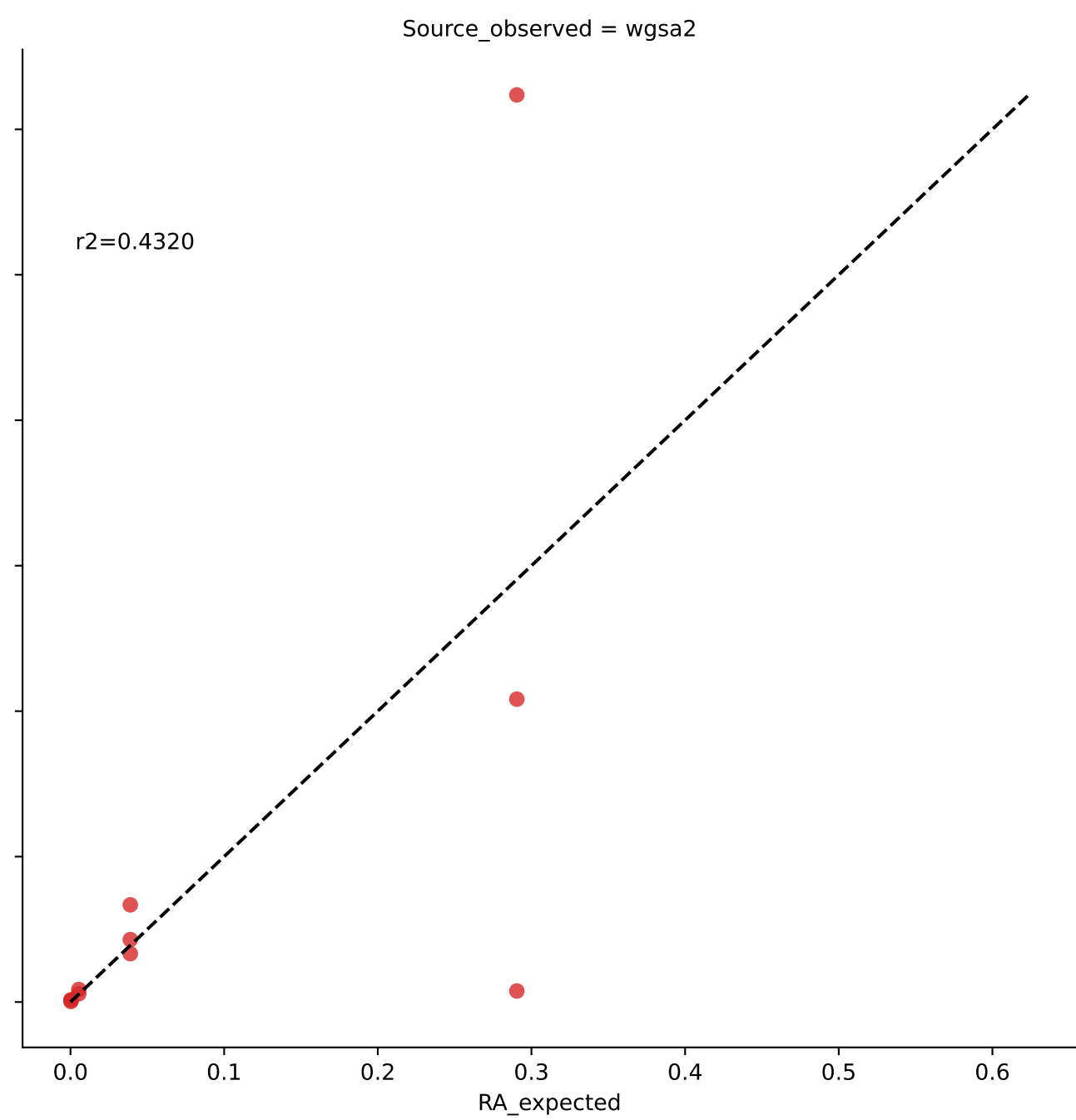
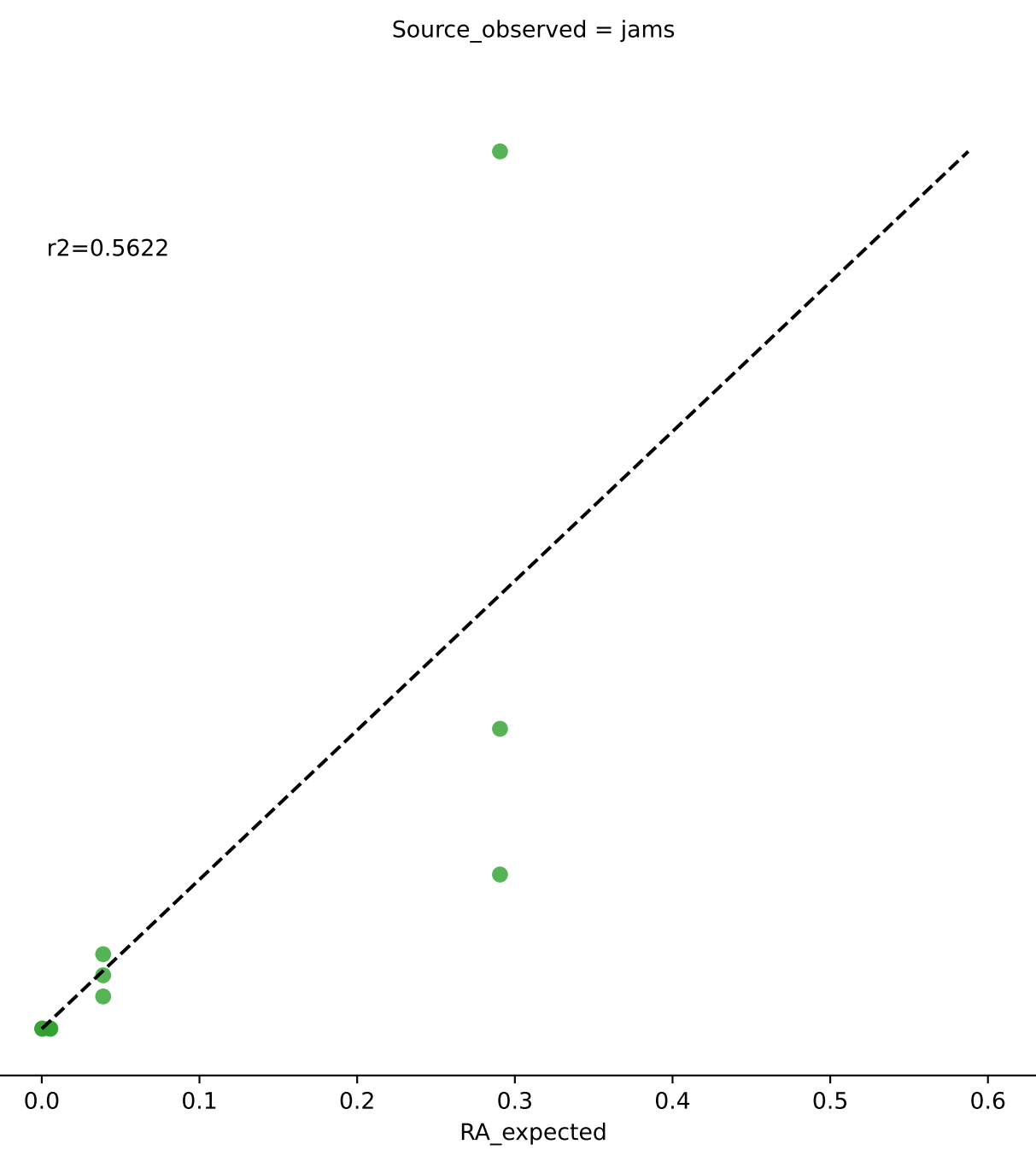
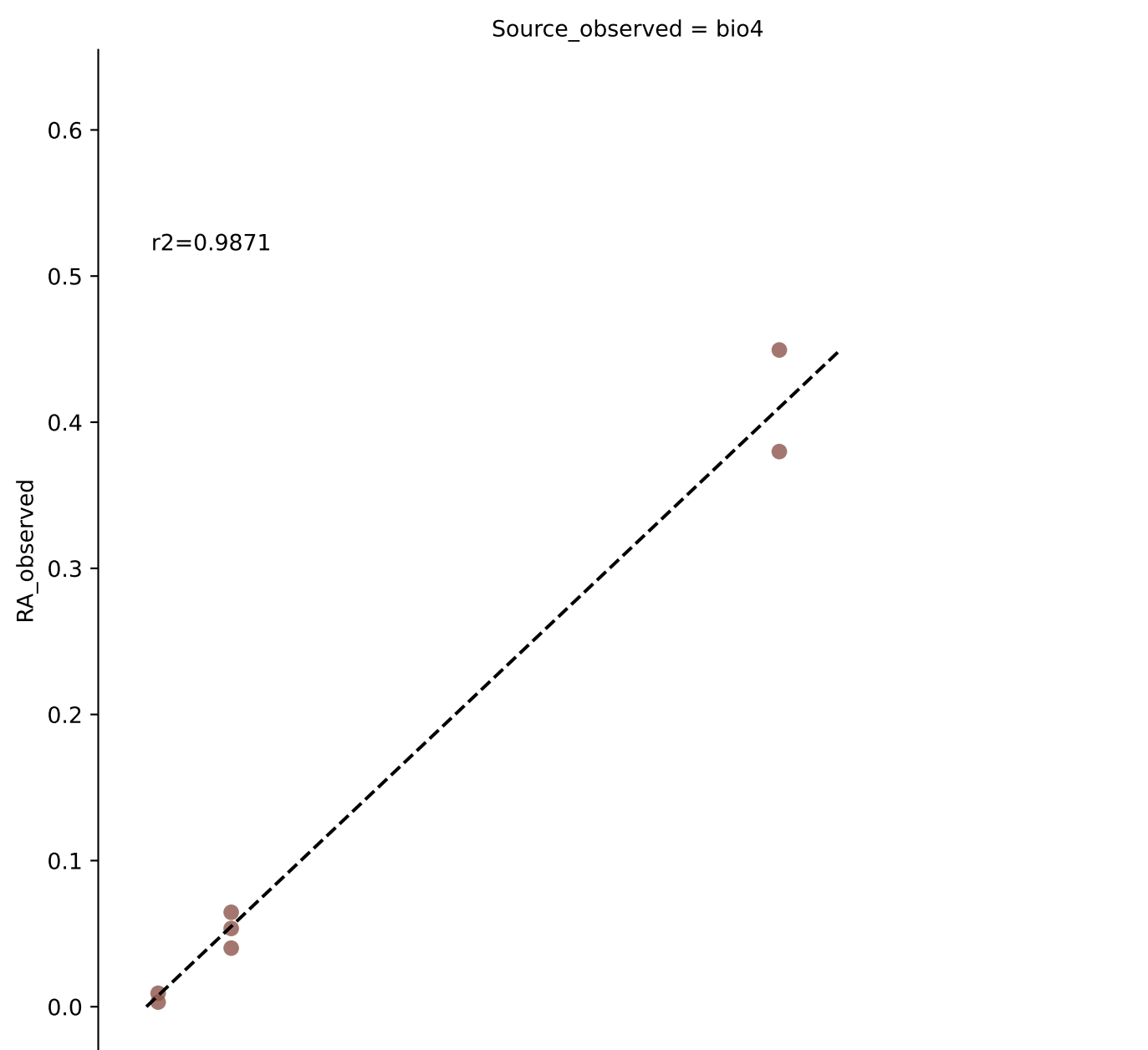


Bivariate Linear Regression for Sample MIX-B in Experiment nist (Genus at filter threshold 1e-05)

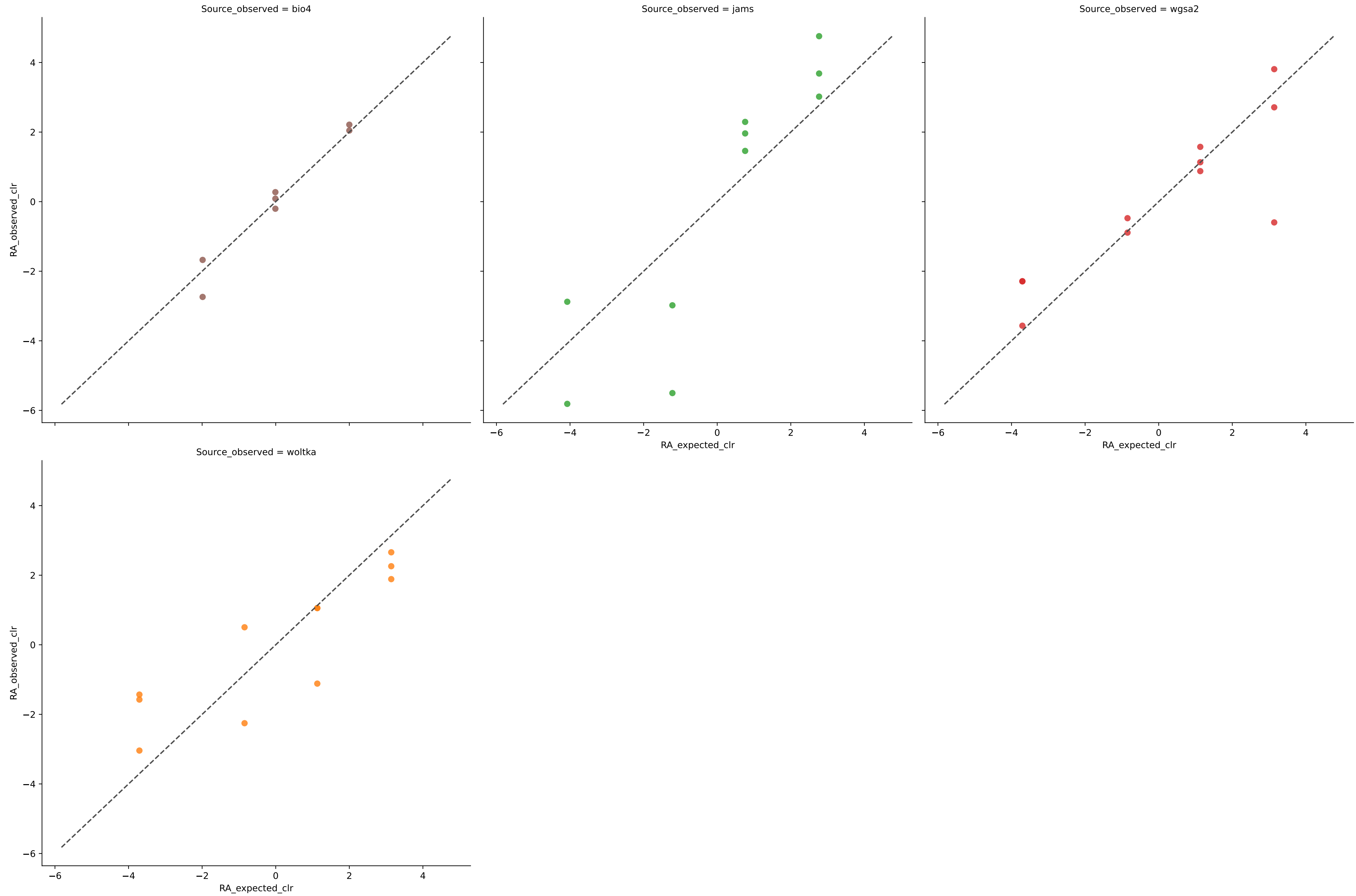


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	8	0.6977	0.0532	4.7646	0.7871	0.0857	100.0000	0.0000
jams	9	0.9604	0.0248	6.3339	0.8885	0.0329	100.0000	0.0000
wgsa2	11	0.9691	0.0167	5.2327	0.9083	0.0241	100.0000	0.0000
woltka	11	0.7965	0.0435	6.1353	0.7607	0.0631	100.0000	0.0000

Bivariate Linear Regression for Sample MIX-C in Experiment nist (Genus at filter threshold 1e-05)

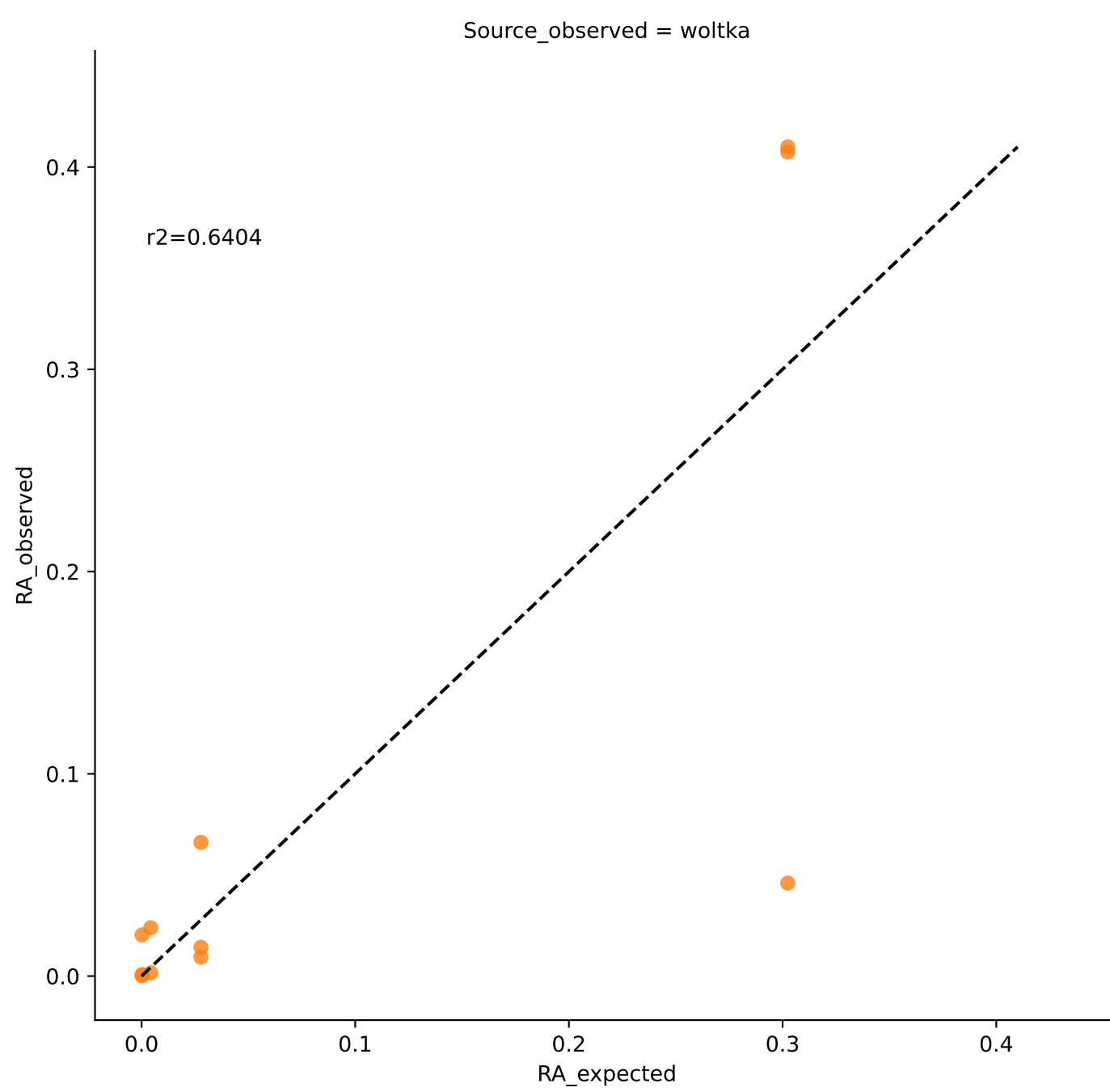
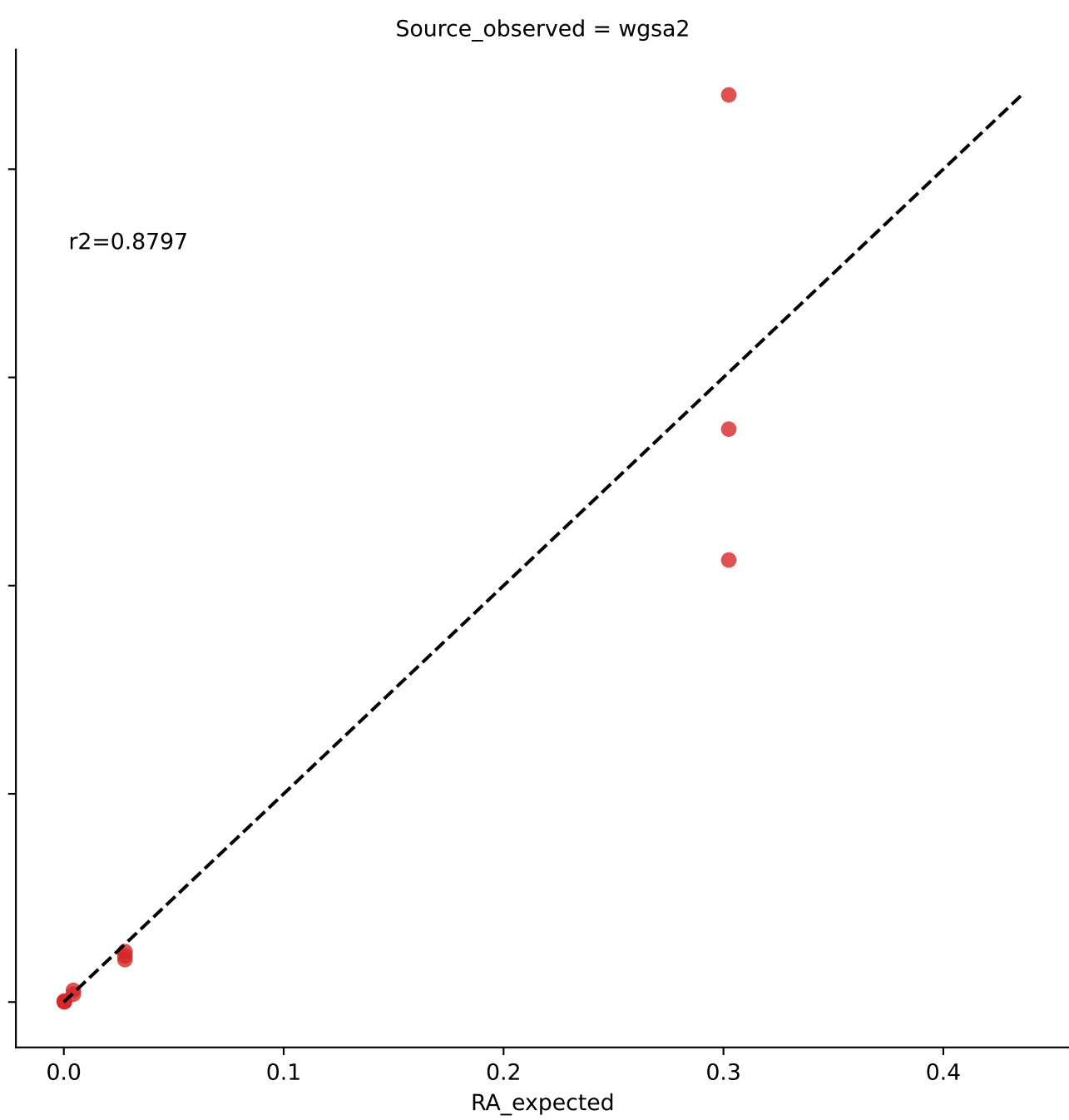
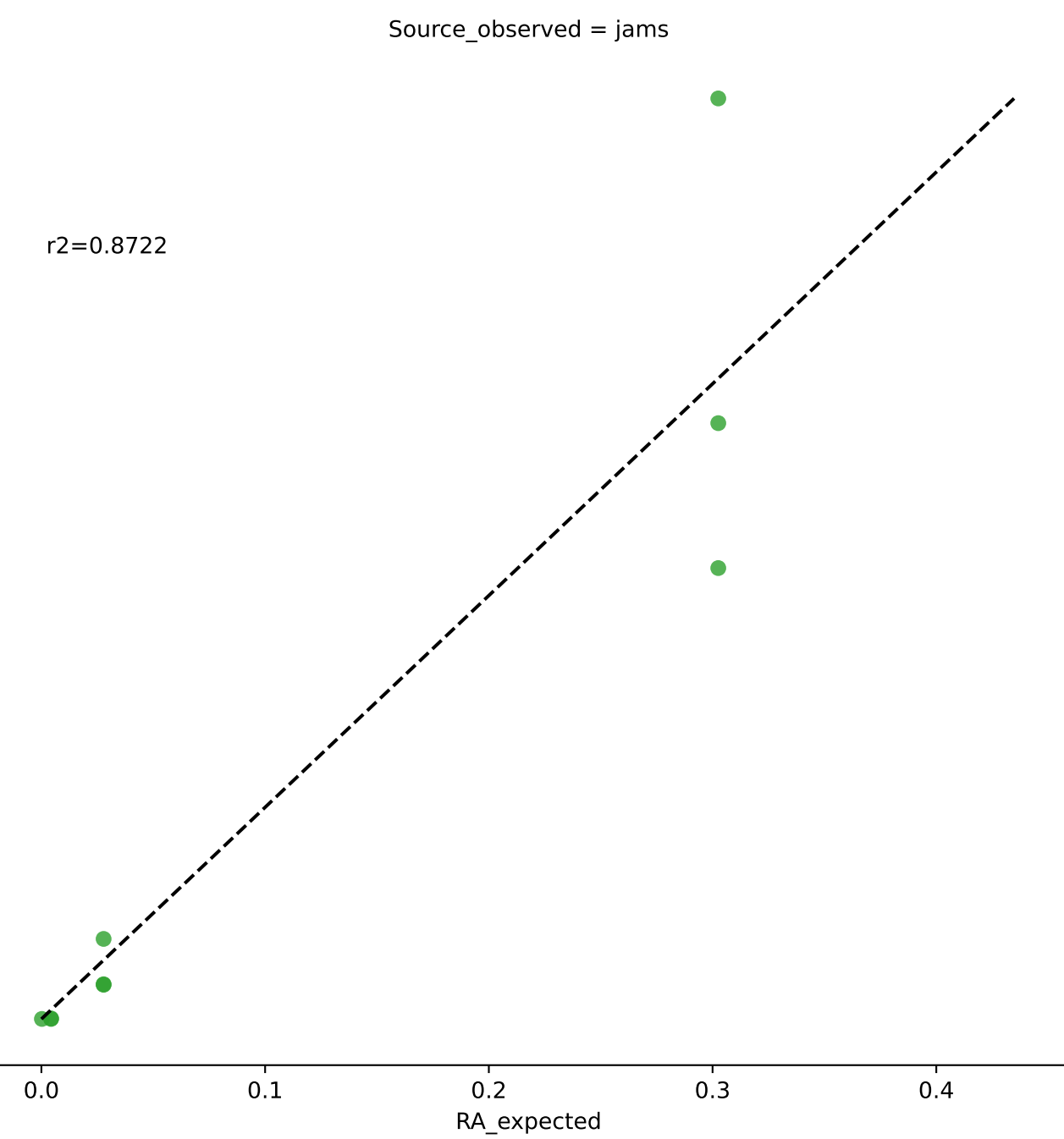
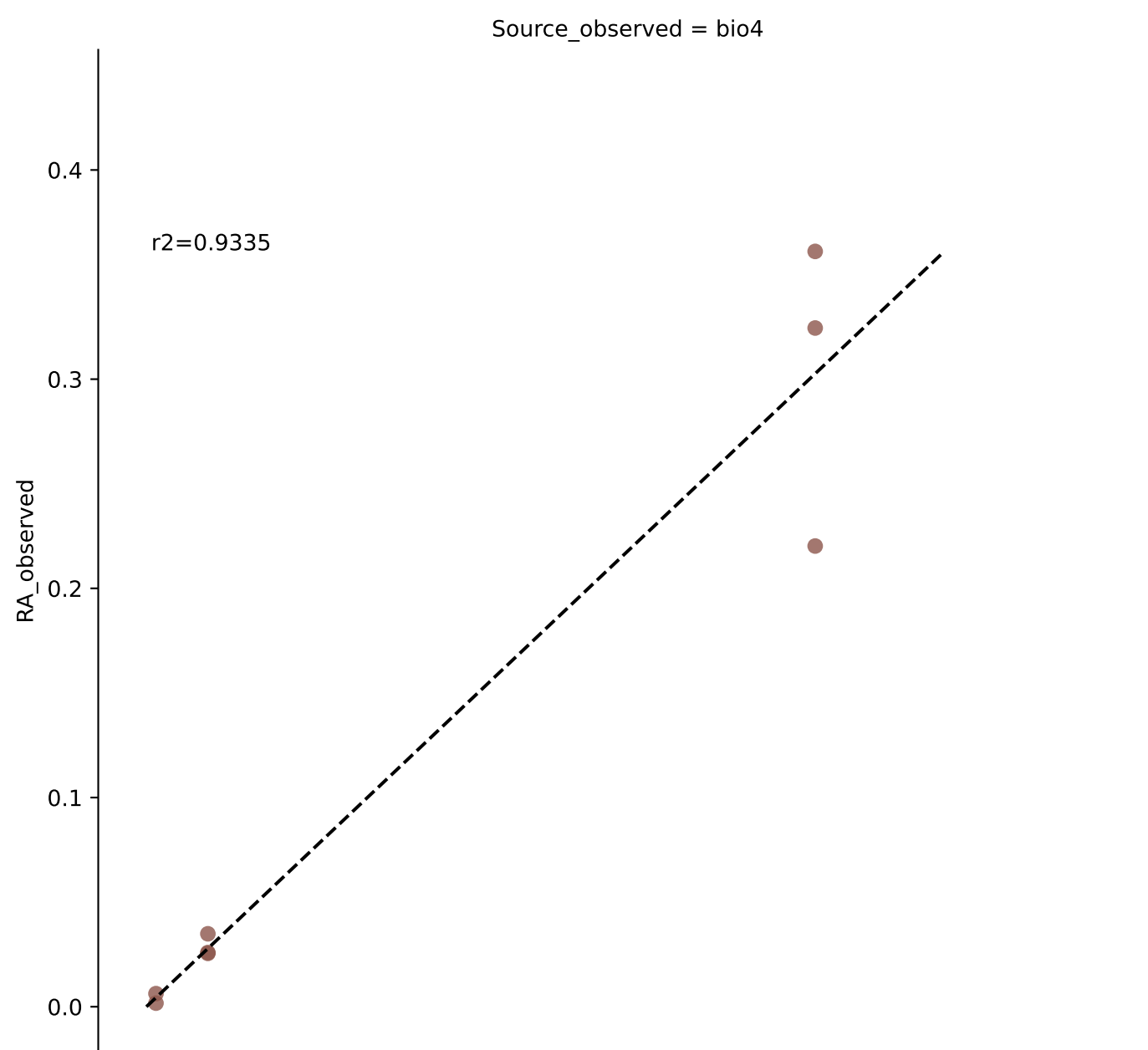


Bivariate Linear Regression for Sample MIX-C in Experiment nist (Genus at filter threshold 1e-05)

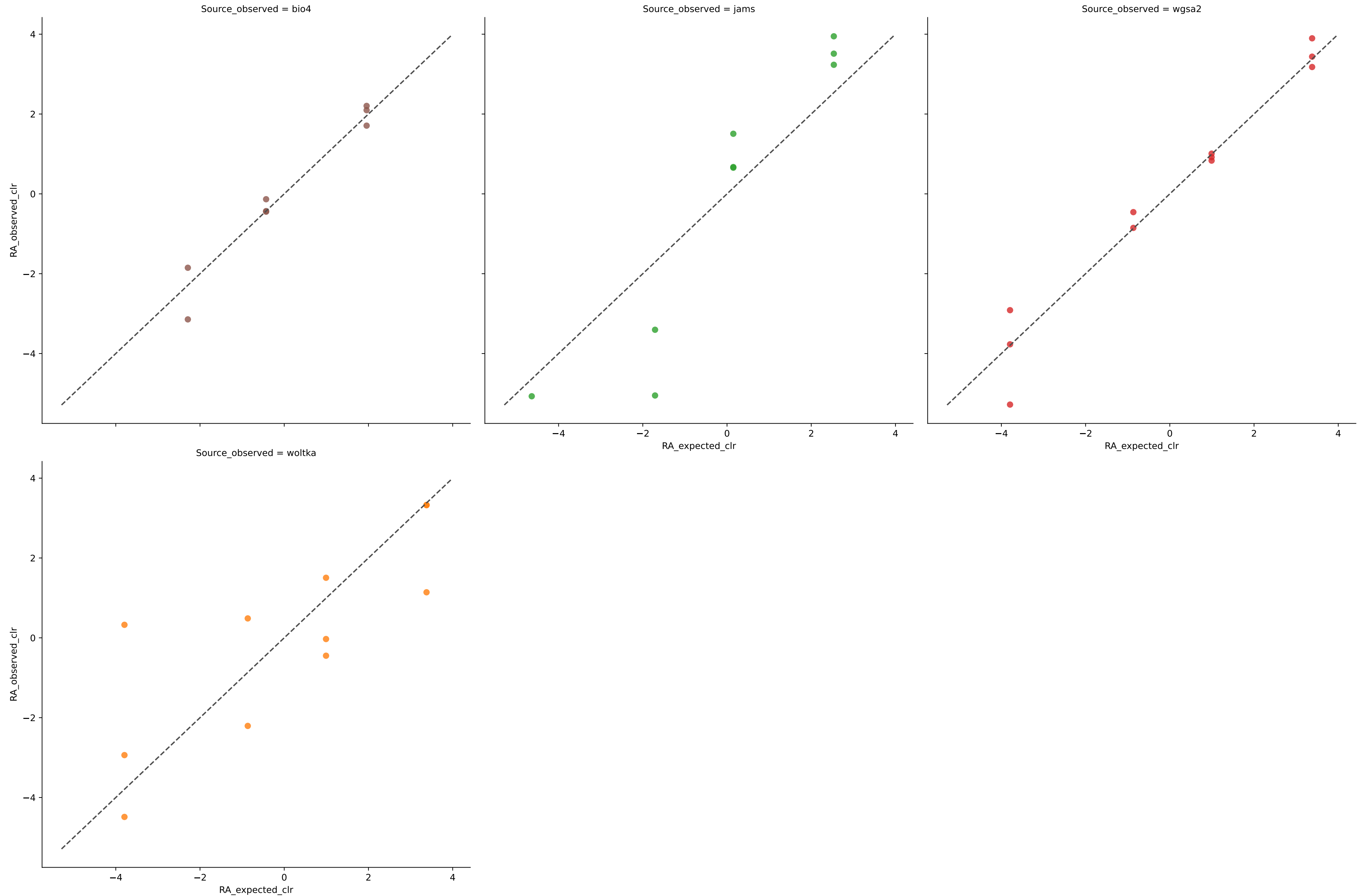


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	7	0.9871	0.0145	0.9149	0.9493	0.0200	100.0000	0.0000
jams	10	0.5622	0.0616	5.9201	0.6920	0.1148	100.0000	0.0000
wgsa2	11	0.4320	0.0674	4.3618	0.6292	0.1344	100.0000	0.0000
woltka	11	0.8372	0.0358	4.6458	0.8029	0.0498	100.0000	0.0000

Bivariate Linear Regression for Sample MIX-D in Experiment nist (Genus at filter threshold 1e-05)

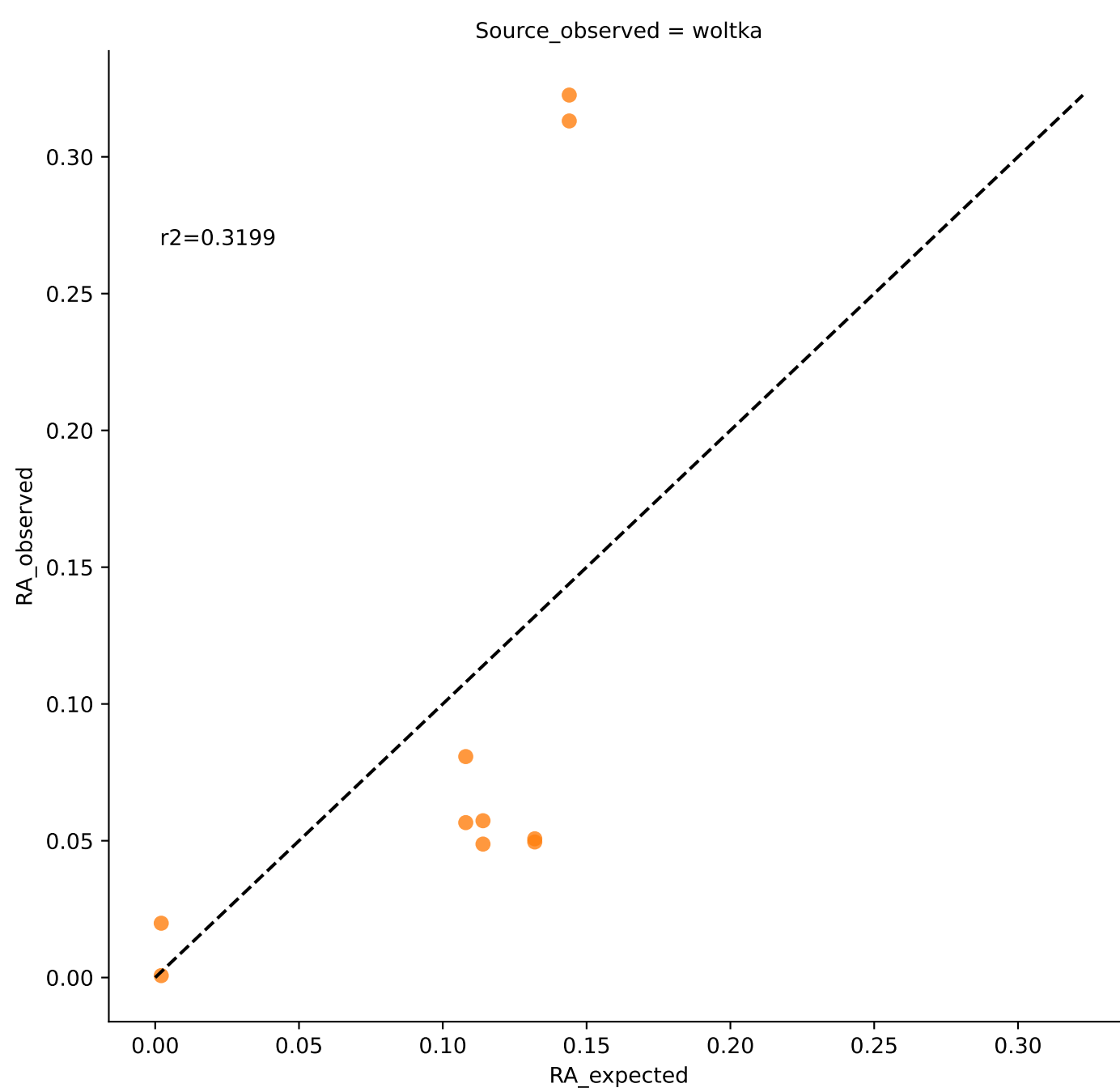
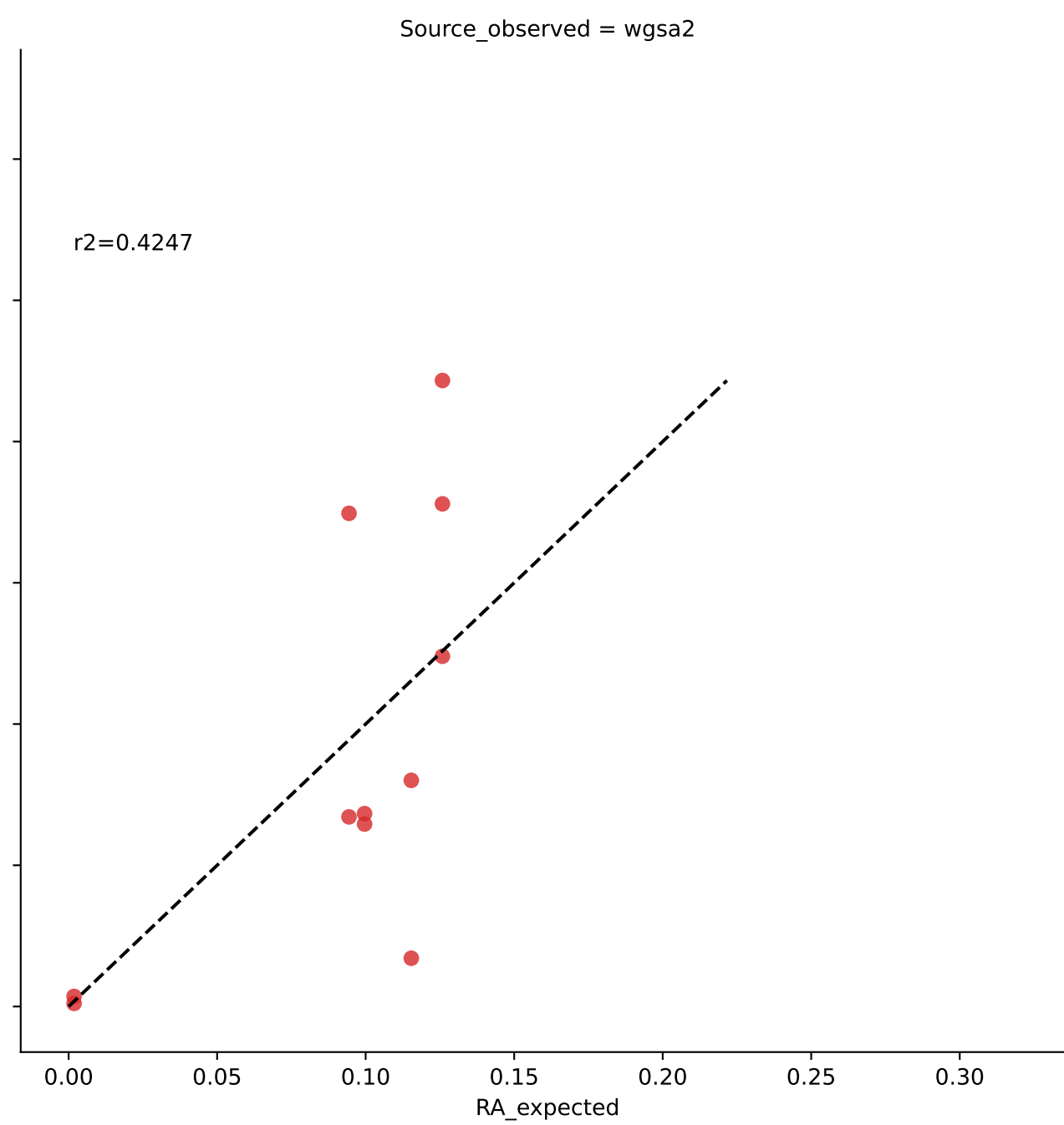
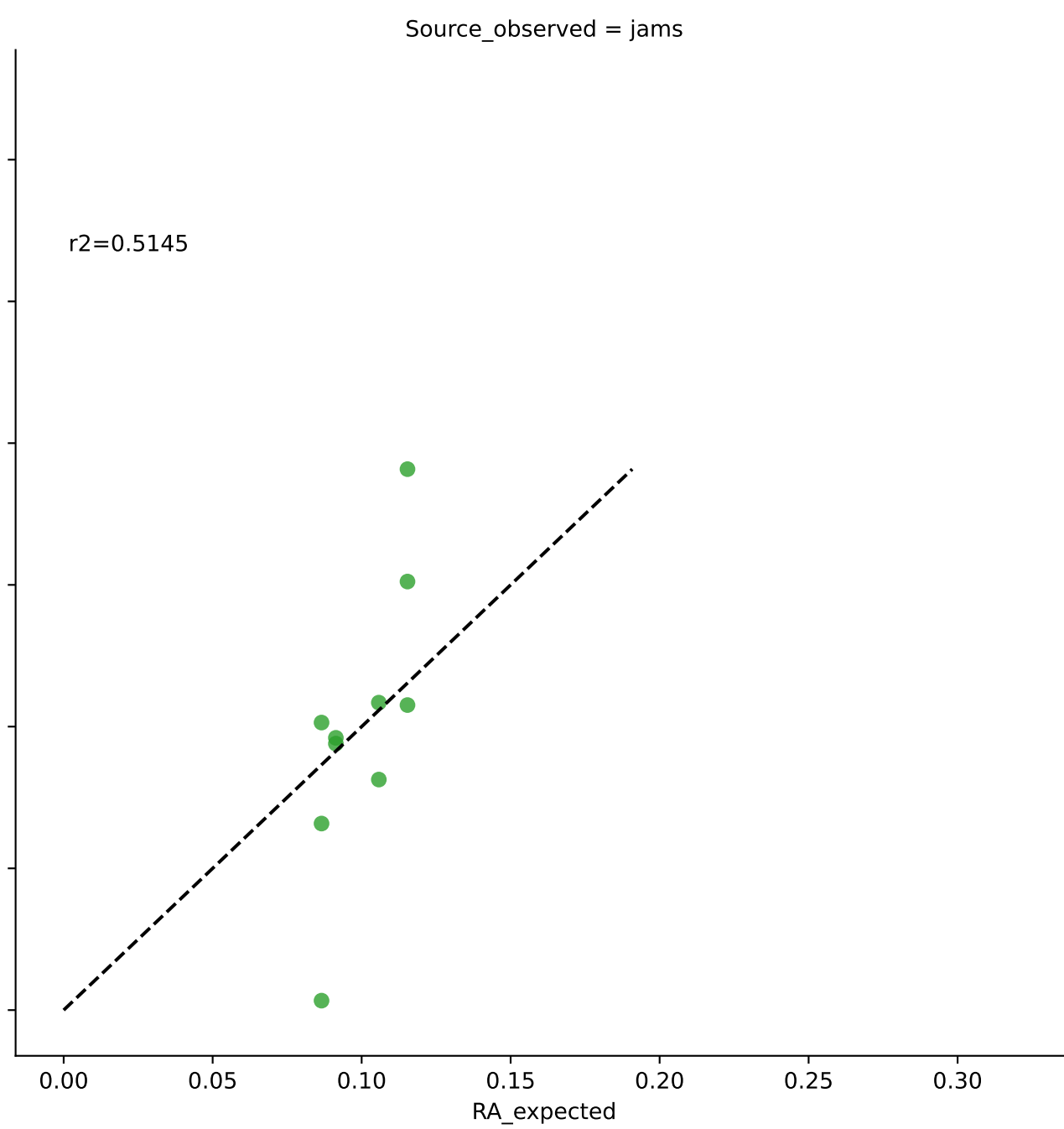
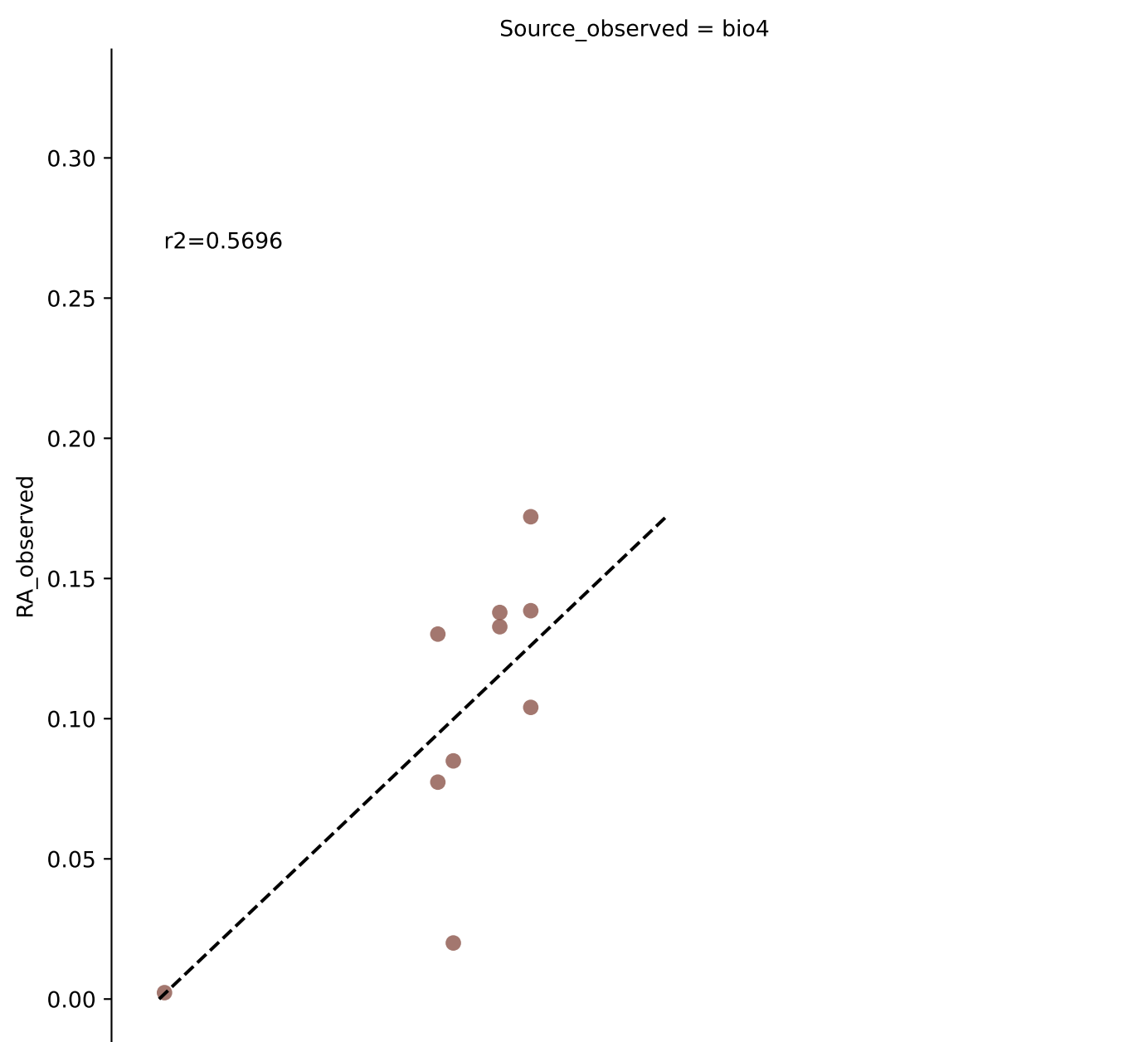


Bivariate Linear Regression for Sample MIX-D in Experiment nist (Genus at filter threshold 1e-05)

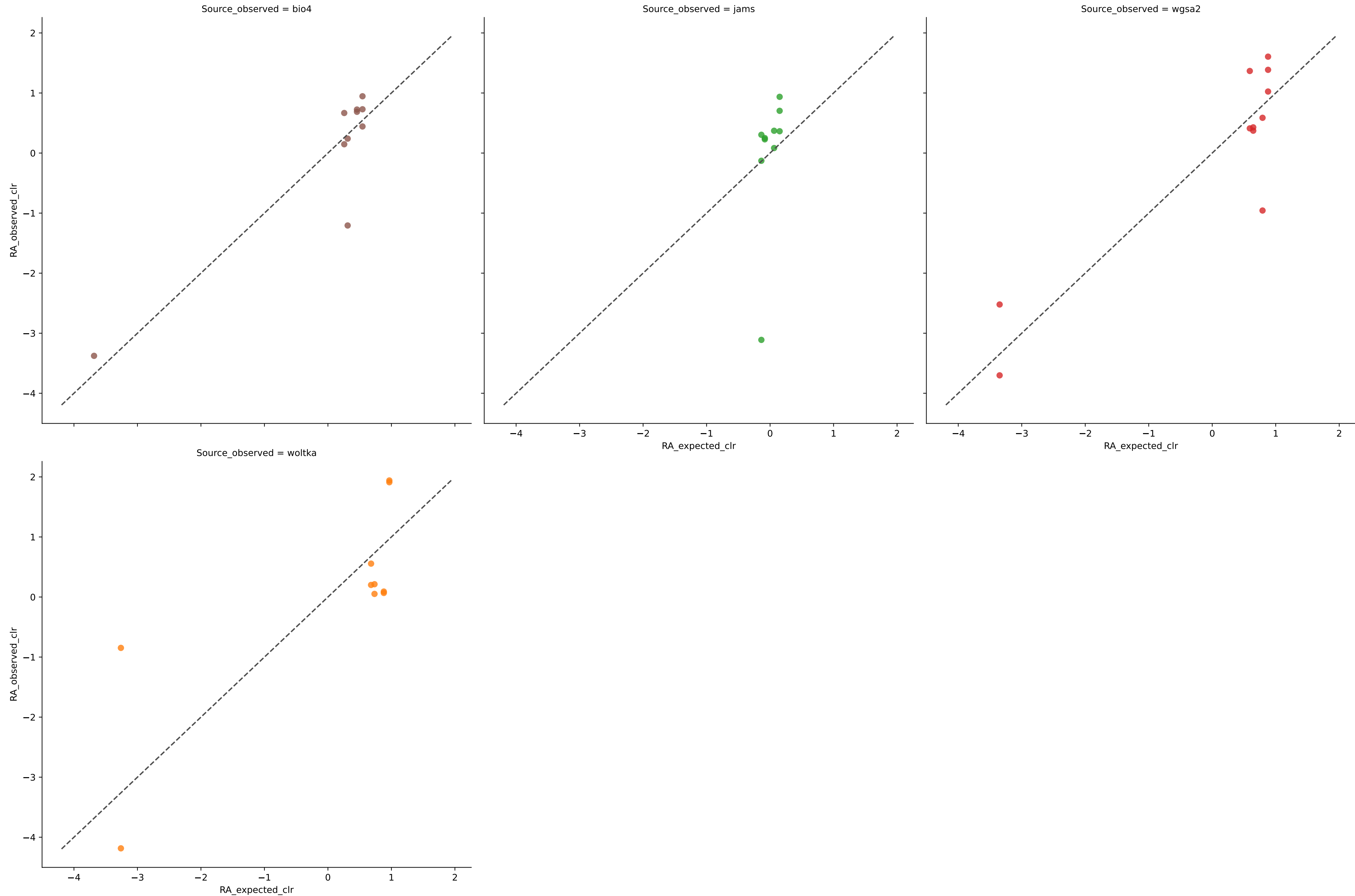


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	8	0.9335	0.0223	1.0739	0.9107	0.0367	100.0000	0.0000
jams	9	0.8722	0.0316	4.4751	0.8578	0.0541	100.0000	0.0000
wgsa2	11	0.8797	0.0245	1.8678	0.8652	0.0493	100.0000	0.0000
woltka	11	0.6404	0.0529	5.4967	0.7088	0.0910	100.0000	0.0000

Bivariate Linear Regression for Sample EG in Experiment nist (Species at filter threshold 1e-05)

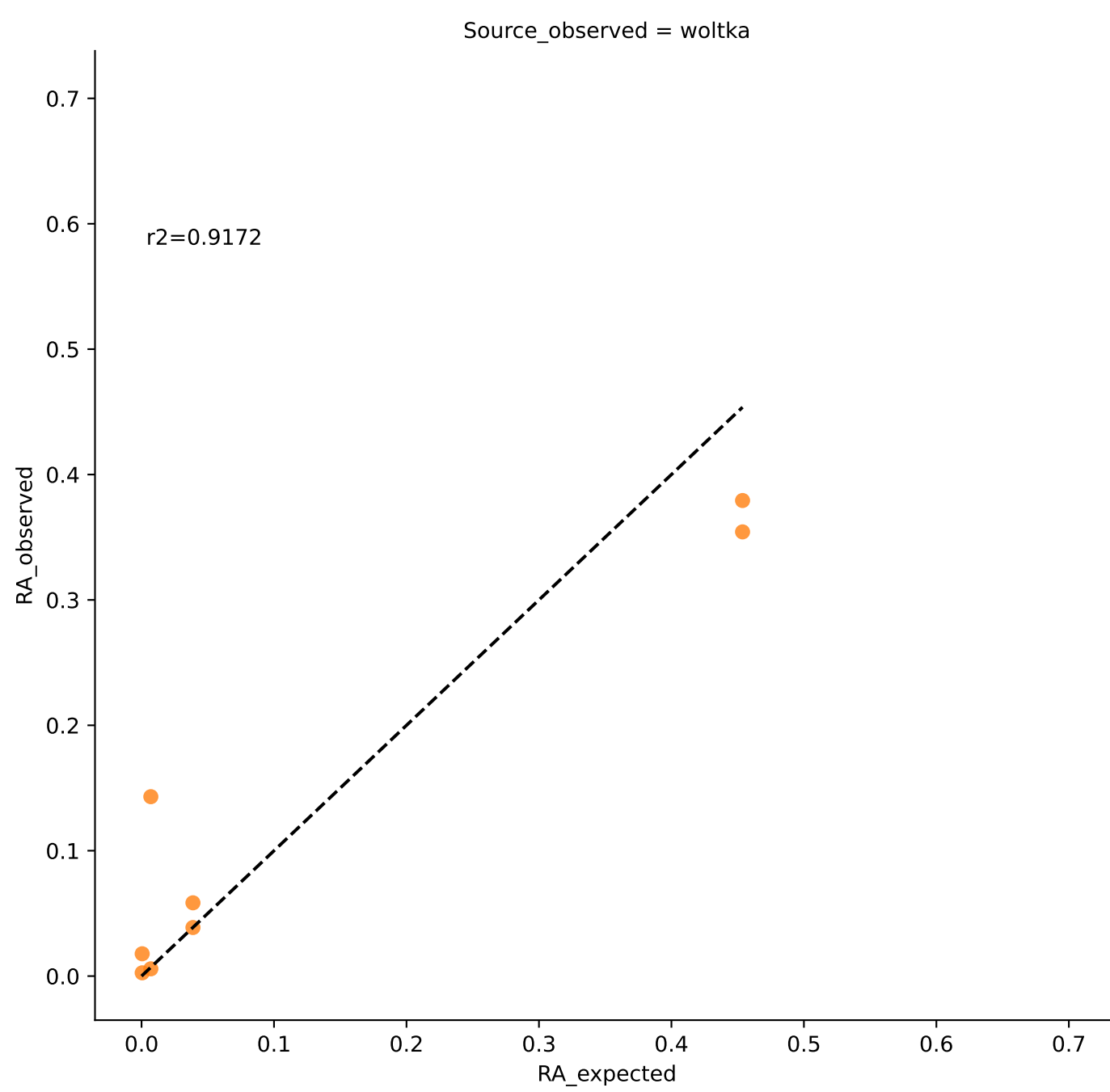
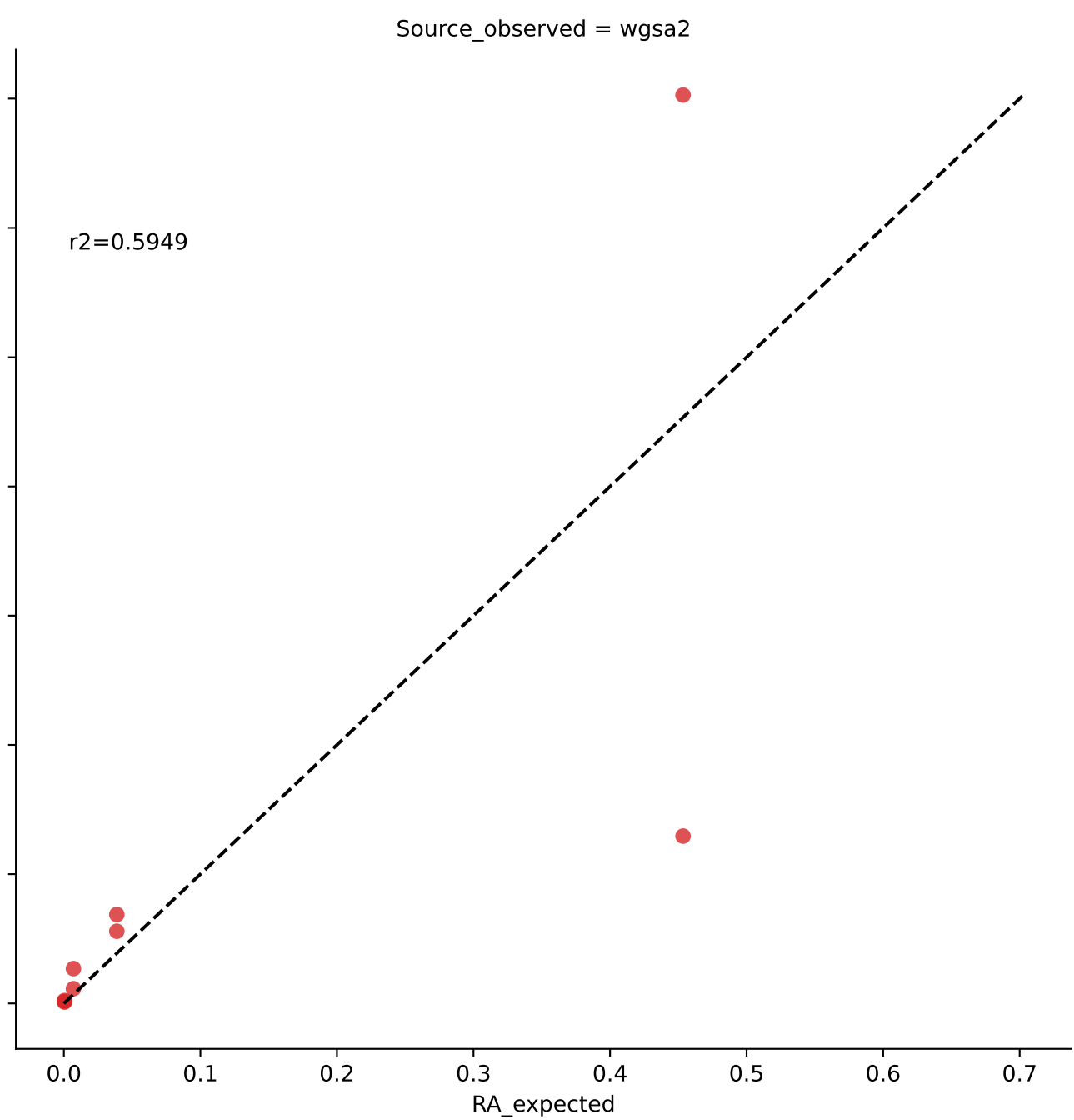
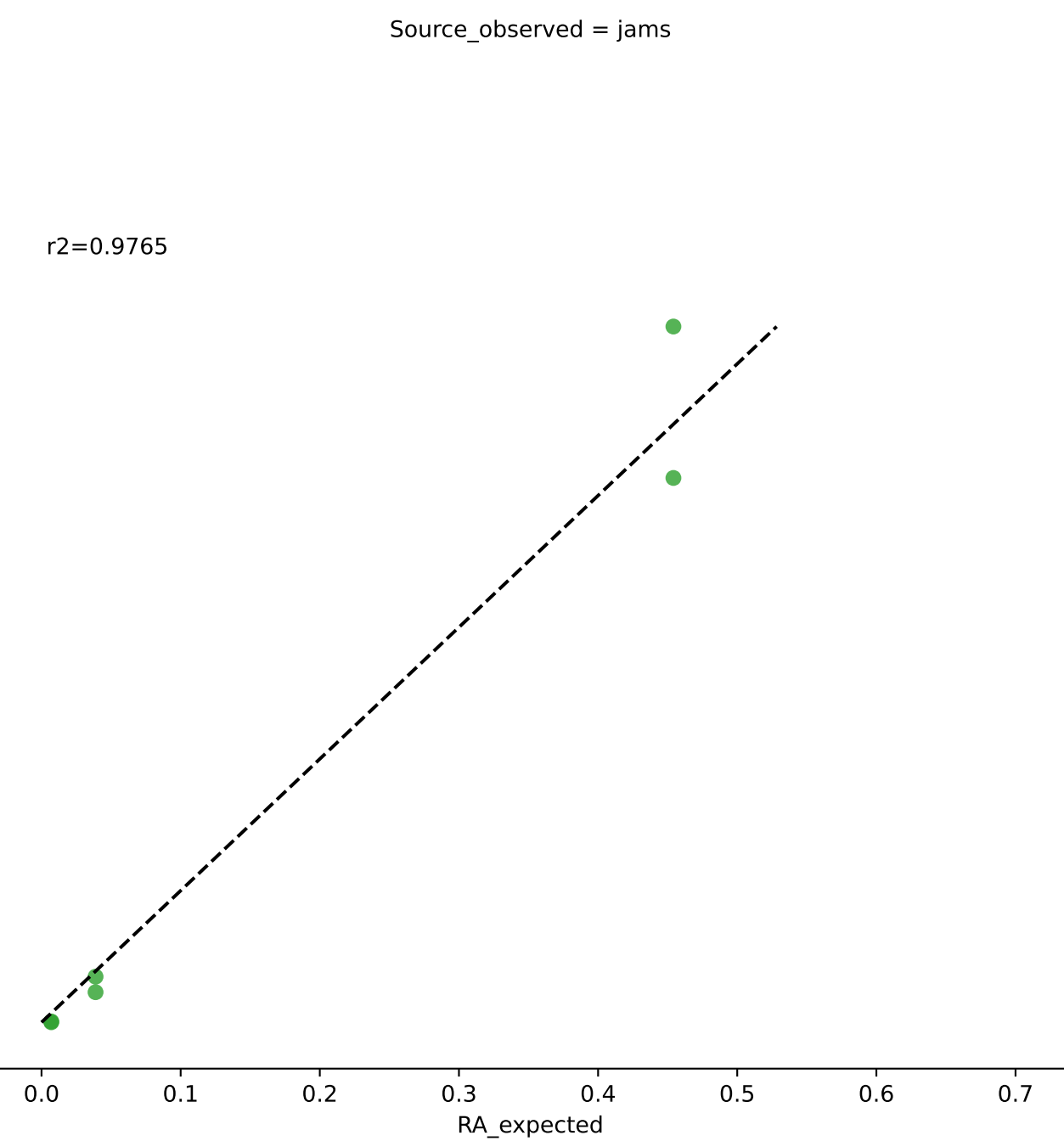
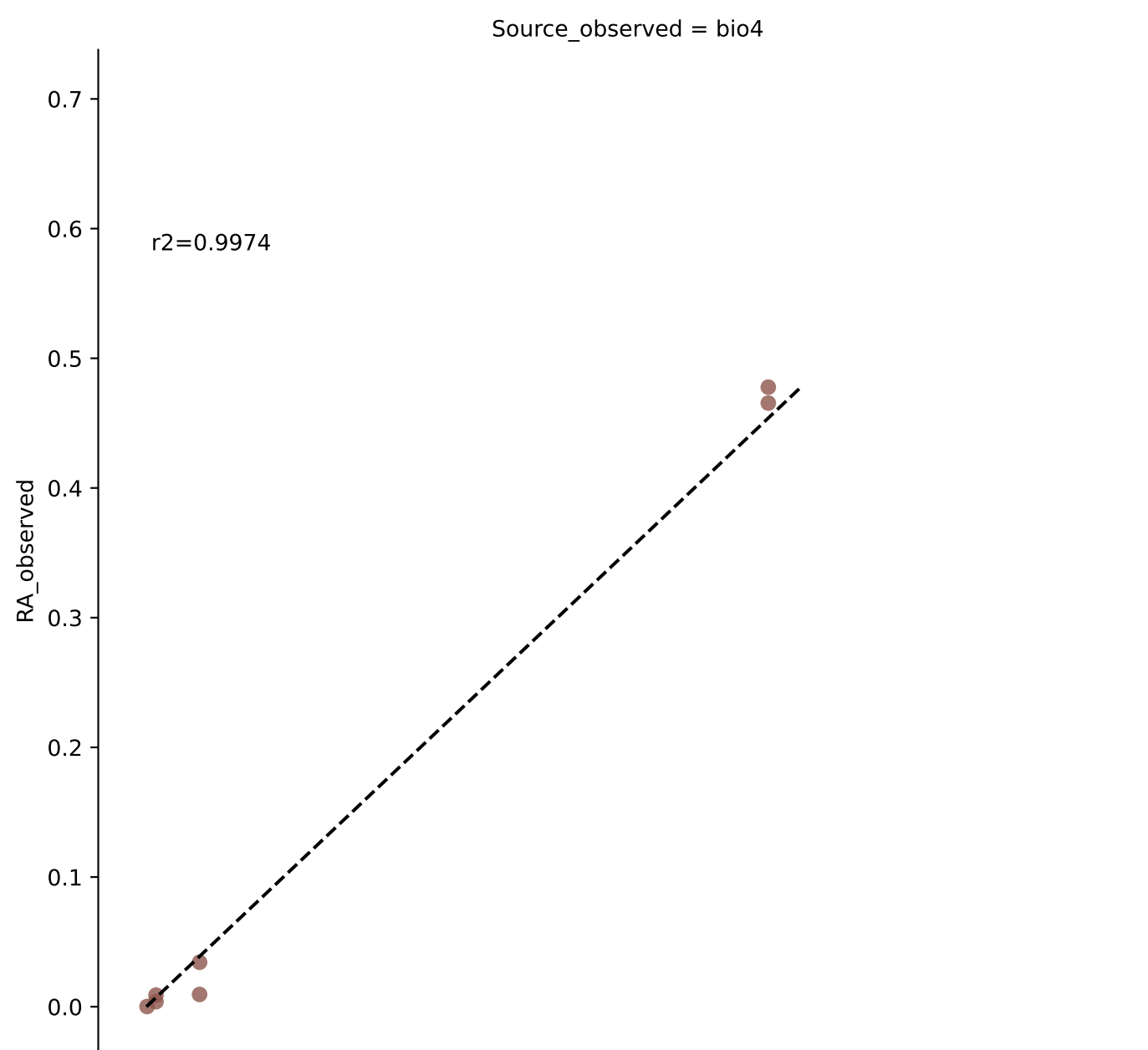


Bivariate Linear Regression for Sample EG in Experiment nist (Species at filter threshold 1e-05)

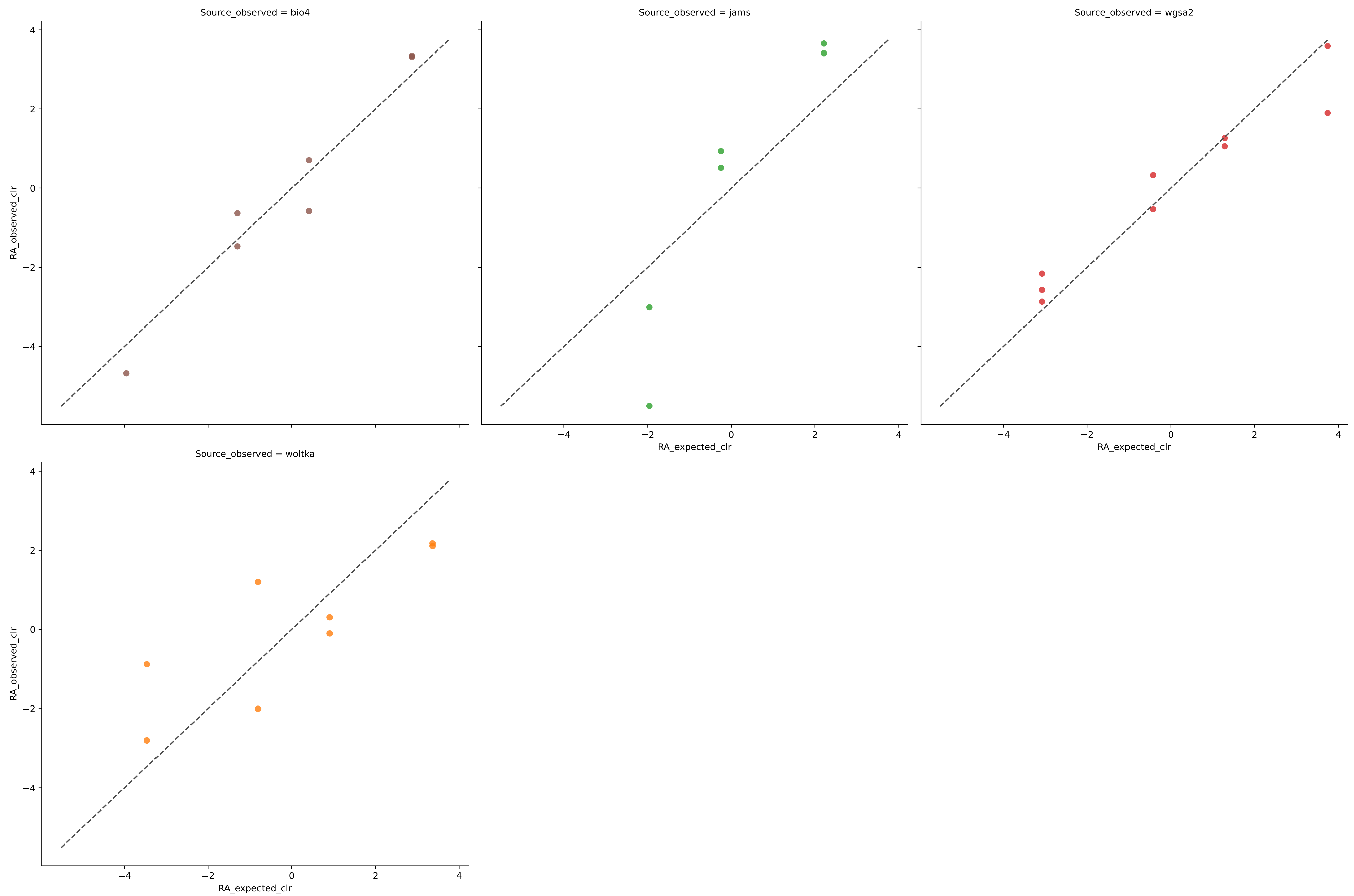


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	10	0.5696	0.0268	1.7058	0.8661	0.0342	100.0000	0.0000
jams	10	0.5145	0.0272	3.2110	0.8638	0.0390	100.0000	0.0000
wgsa2	11	0.4247	0.0418	2.3389	0.7702	0.0541	100.0000	0.0000
woltka	10	0.3199	0.0731	3.2836	0.6345	0.0922	100.0000	0.0000

Bivariate Linear Regression for Sample MIX-A in Experiment nist (Species at filter threshold 1e-05)

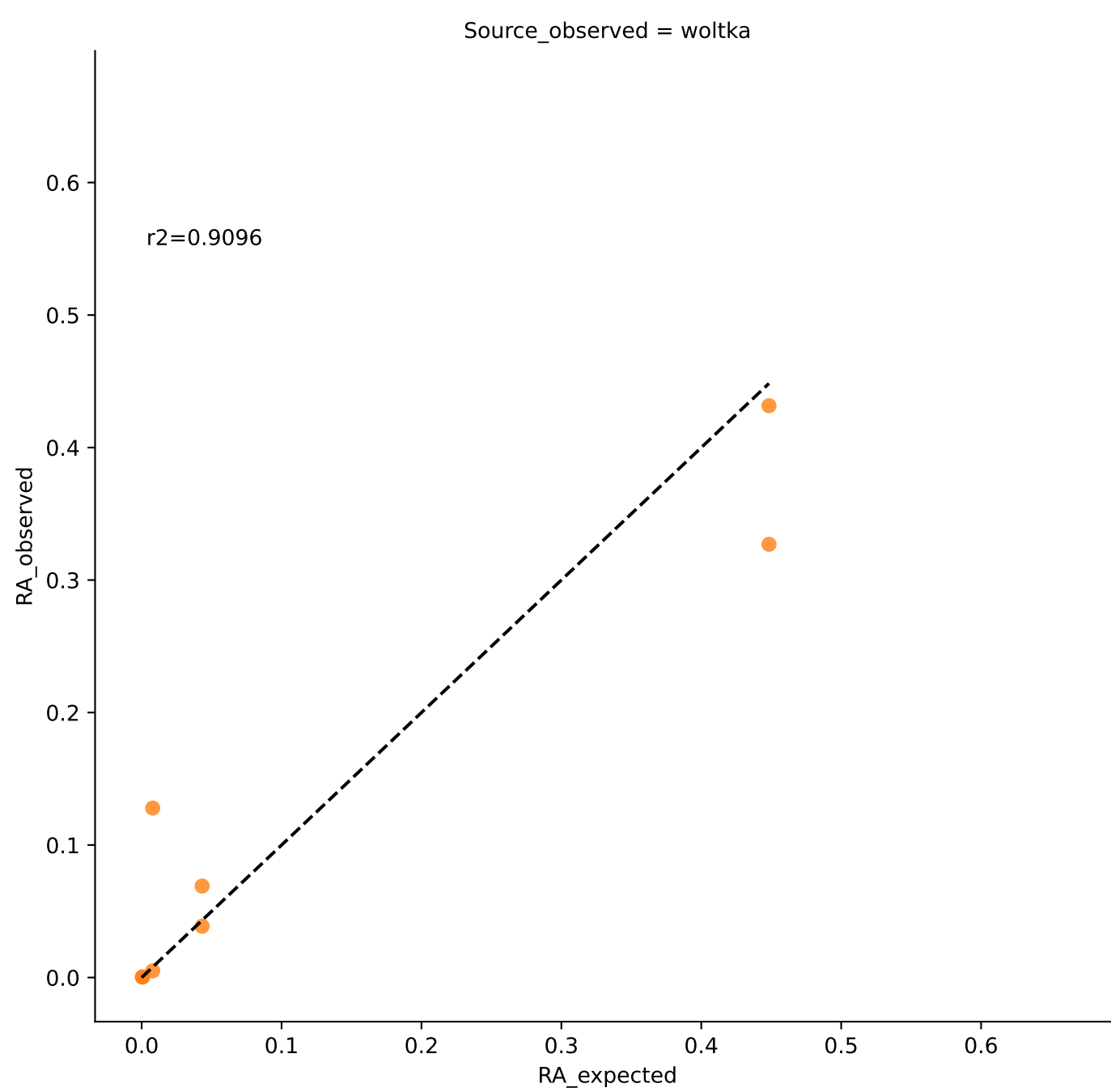
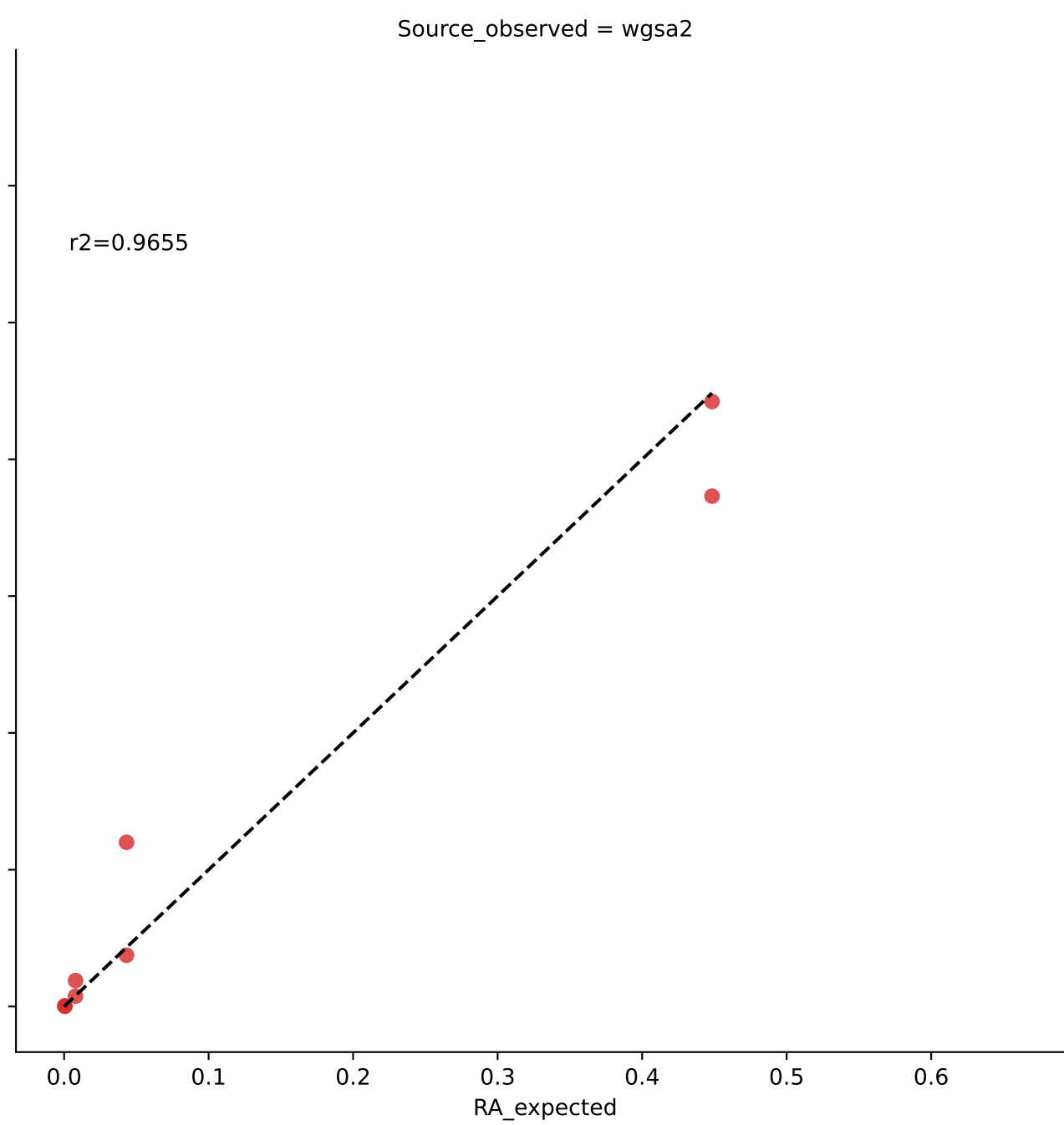
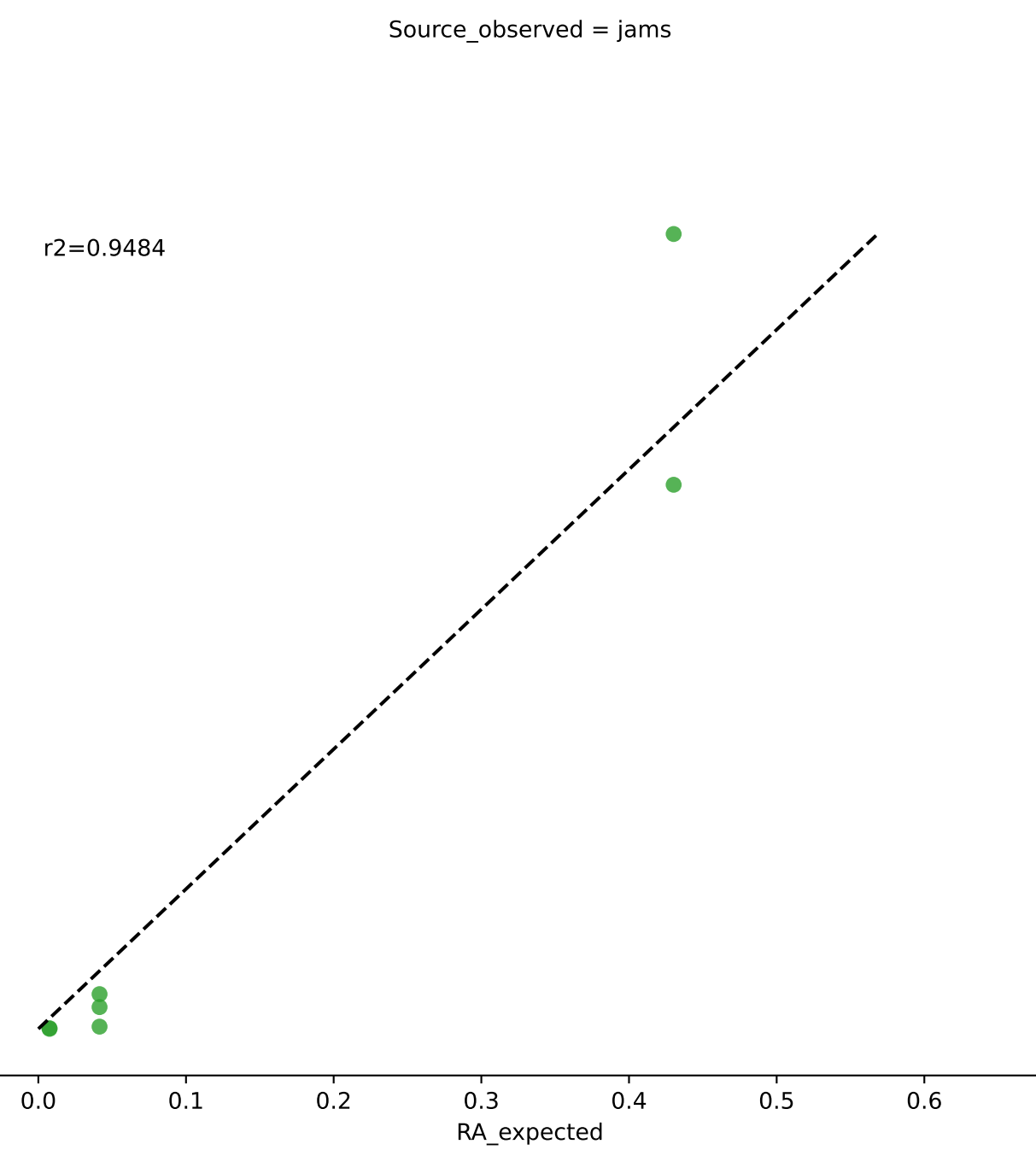
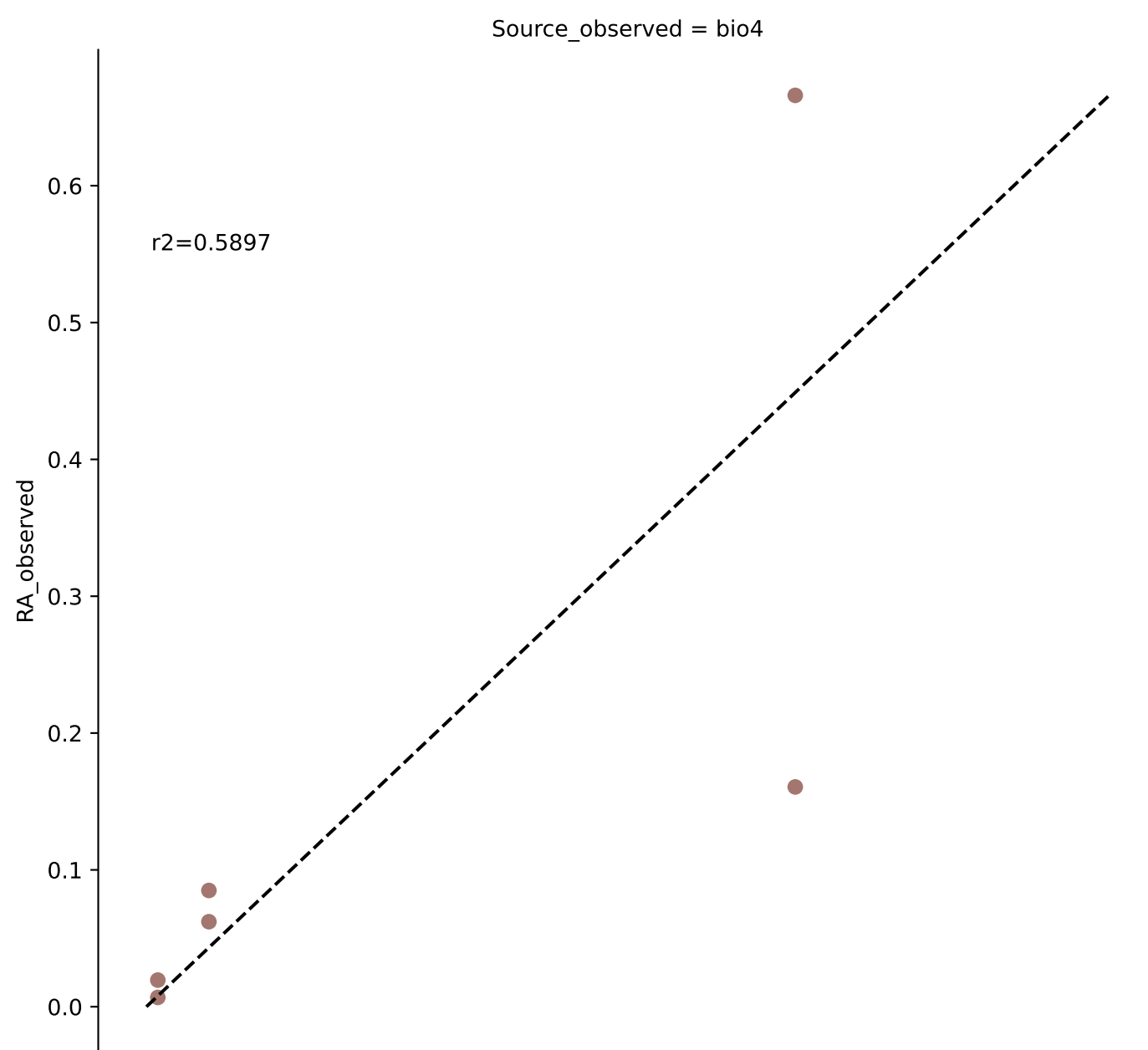


Bivariate Linear Regression for Sample MIX-A in Experiment nist (Species at filter threshold 1e-05)

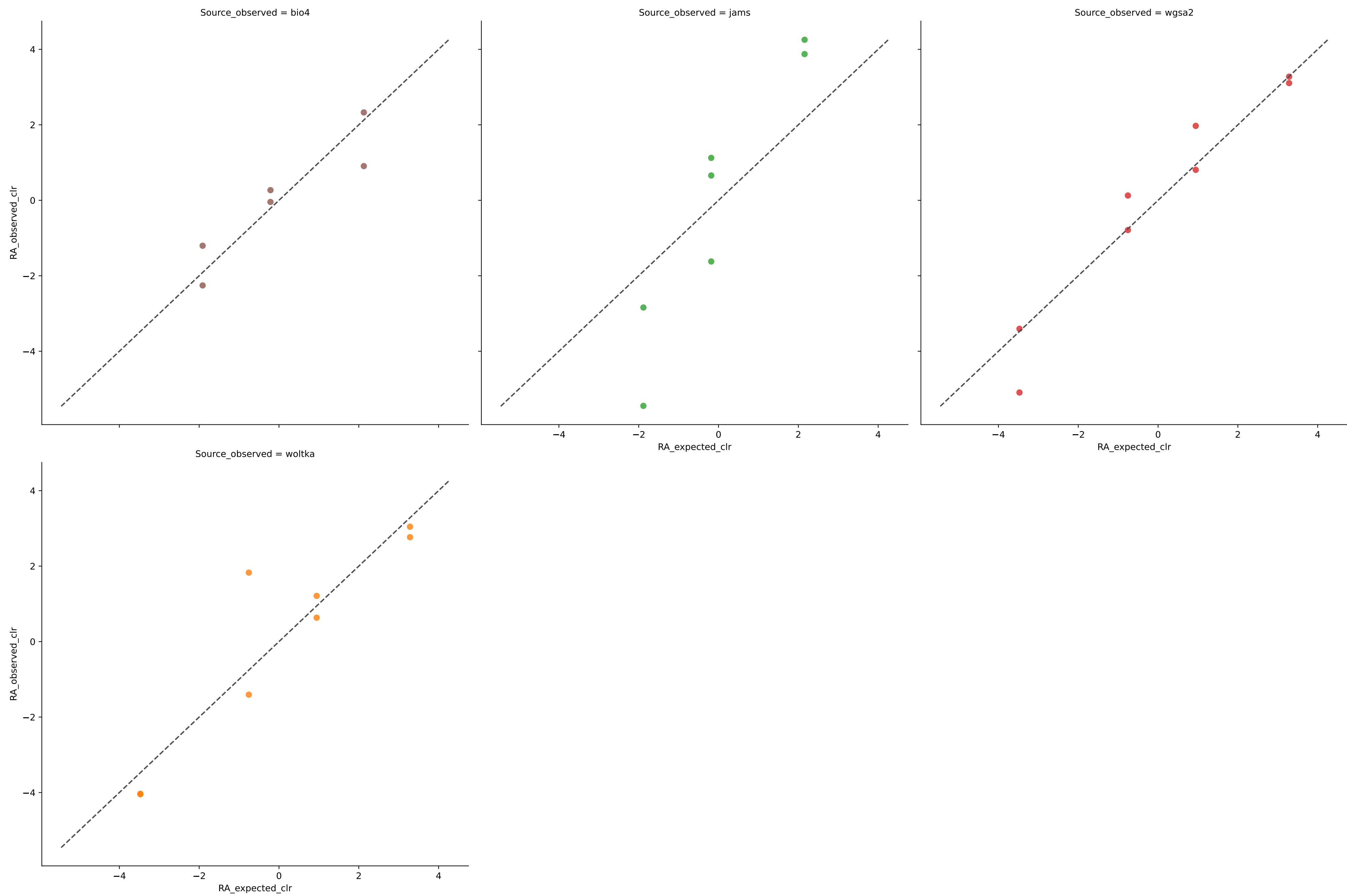


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	7	0.9974	0.0107	1.5752	0.9626	0.0151	100.0000	0.0000
jams	6	0.9765	0.0247	4.3745	0.9258	0.0354	100.0000	0.0000
wgsa2	9	0.5949	0.0720	2.2860	0.6760	0.1369	100.0000	0.0000
woltka	8	0.9172	0.0438	4.1130	0.8249	0.0658	100.0000	0.0000

Bivariate Linear Regression for Sample MIX-B in Experiment nist (Species at filter threshold 1e-05)

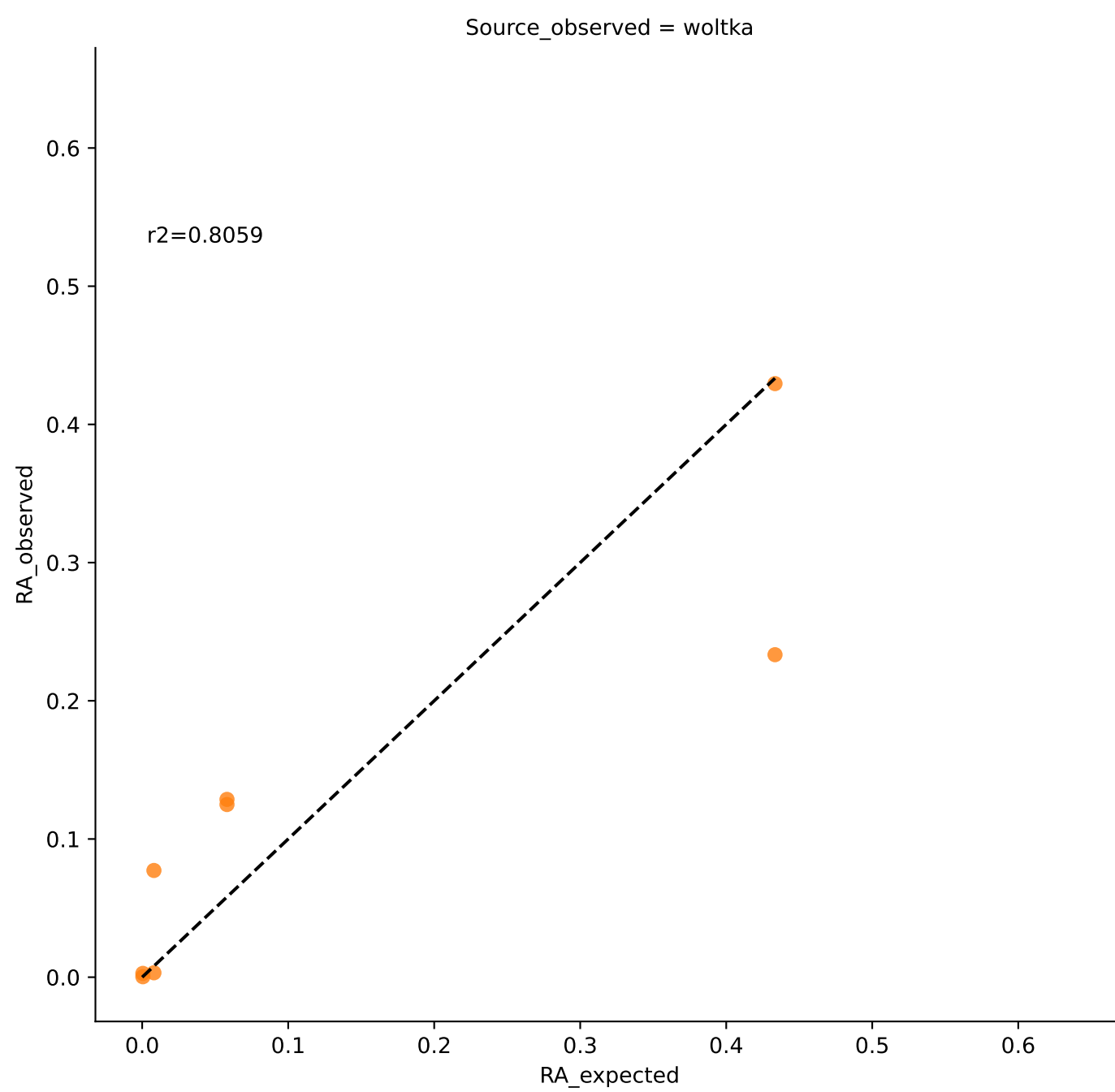
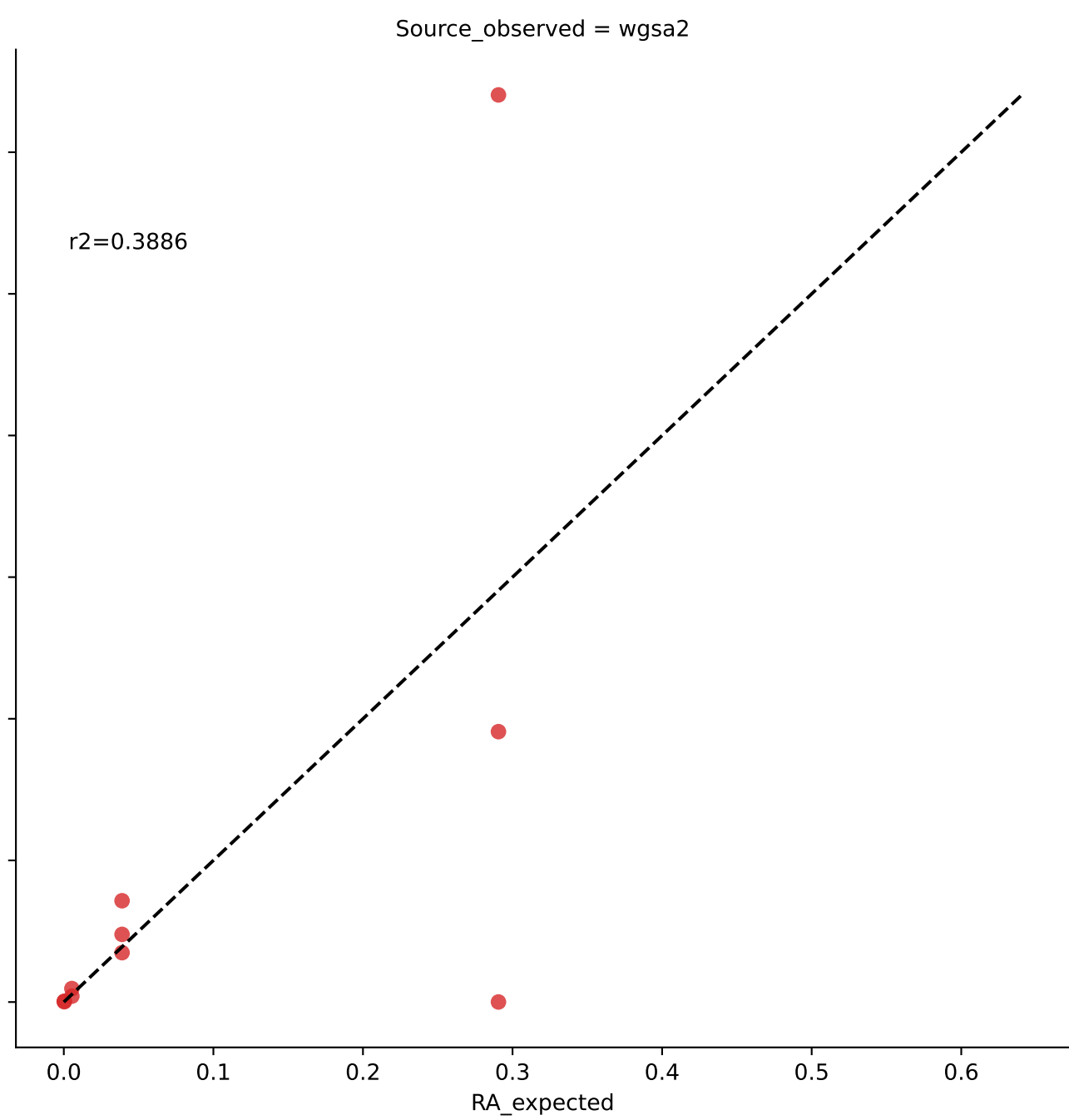
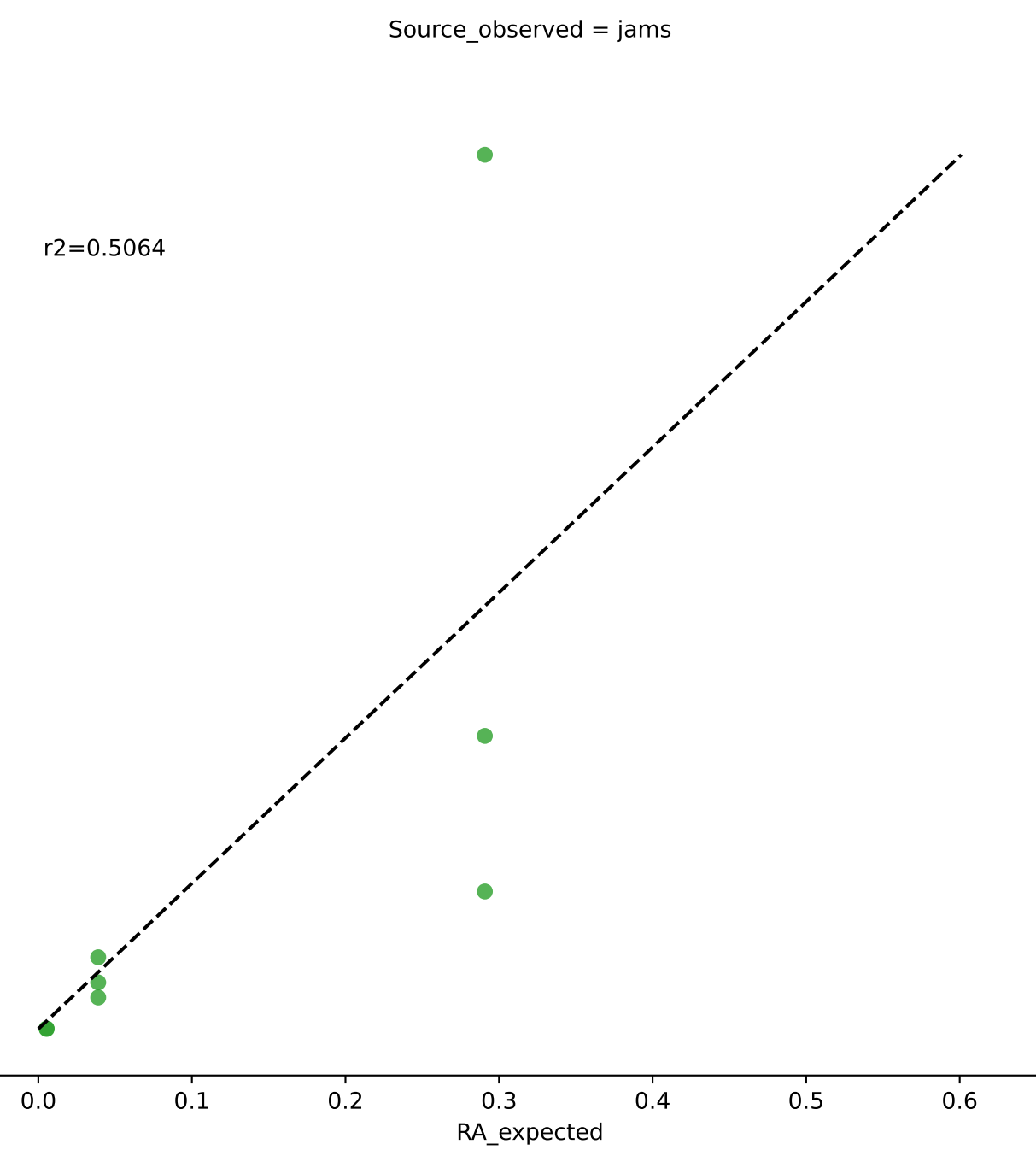
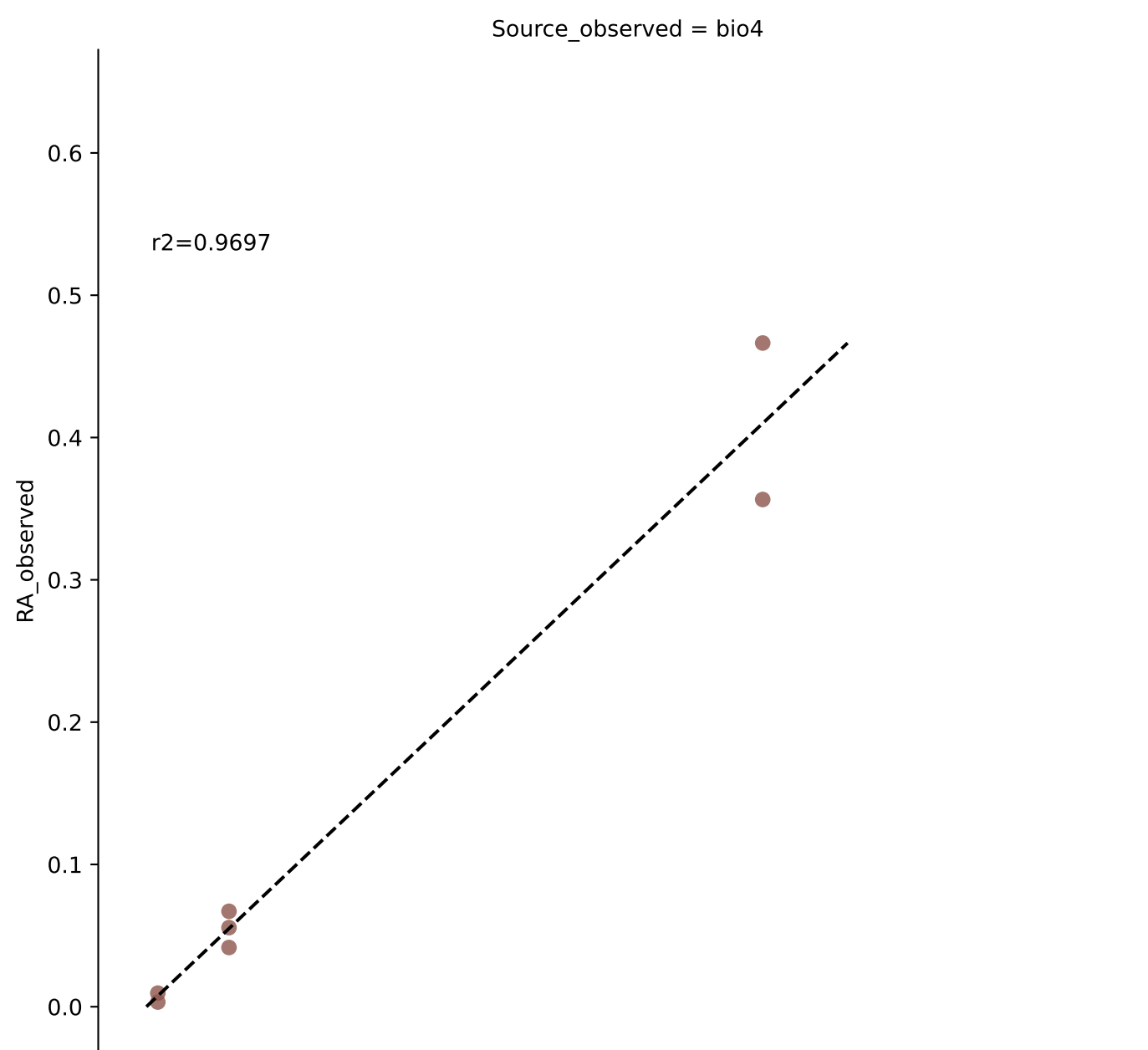


Bivariate Linear Regression for Sample MIX-B in Experiment nist (Species at filter threshold 1e-05)

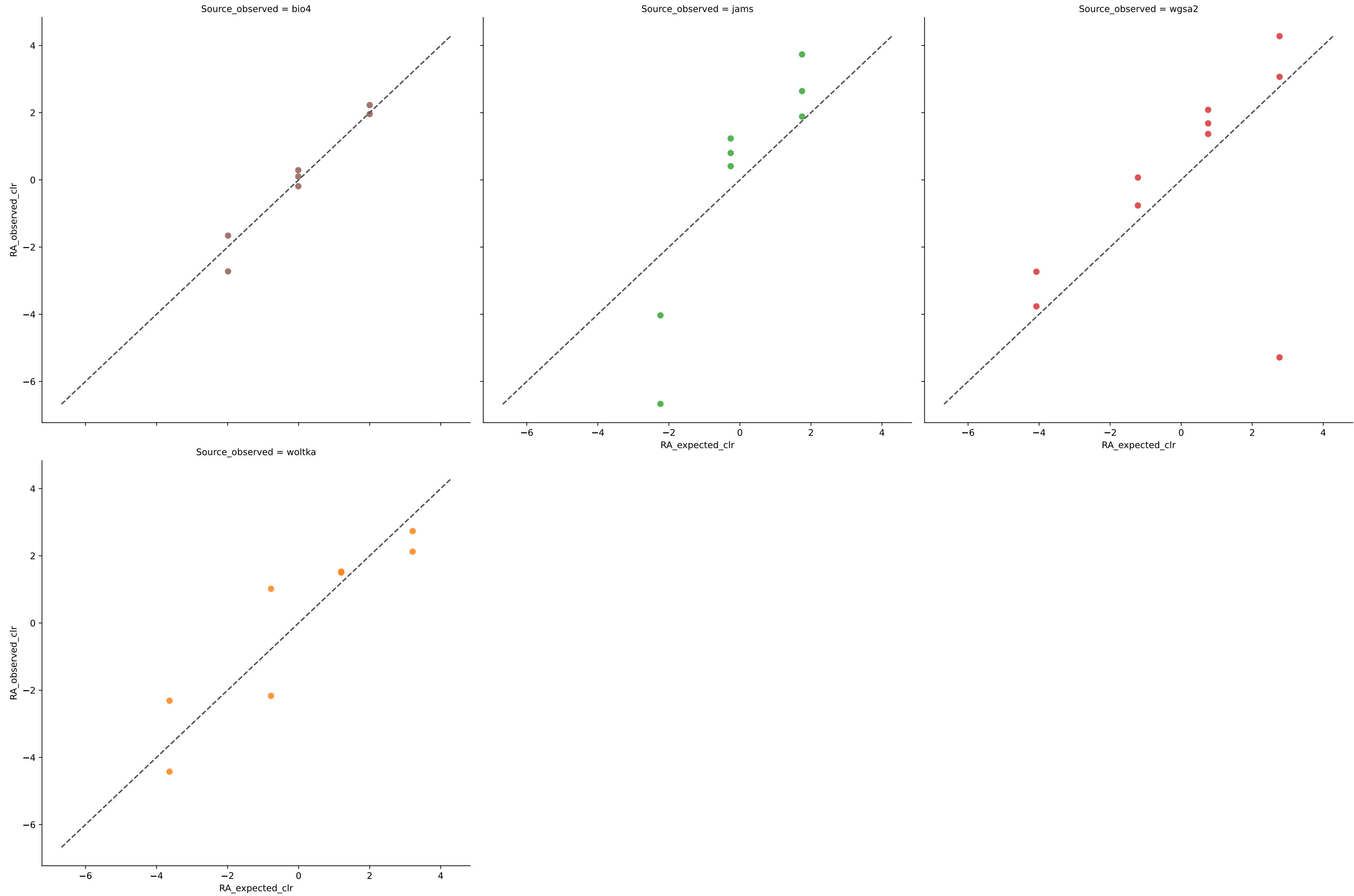


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	6	0.5897	0.0964	1.5530	0.7107	0.1486	100.0000	0.0000
jams	7	0.9484	0.0395	5.0450	0.8619	0.0578	100.0000	0.0000
wgsa2	8	0.9655	0.0220	2.1248	0.9120	0.0384	100.0000	0.0000
woltka	8	0.9096	0.0365	2.8687	0.8542	0.0614	100.0000	0.0000

Bivariate Linear Regression for Sample MIX-C in Experiment nist (Species at filter threshold 1e-05)

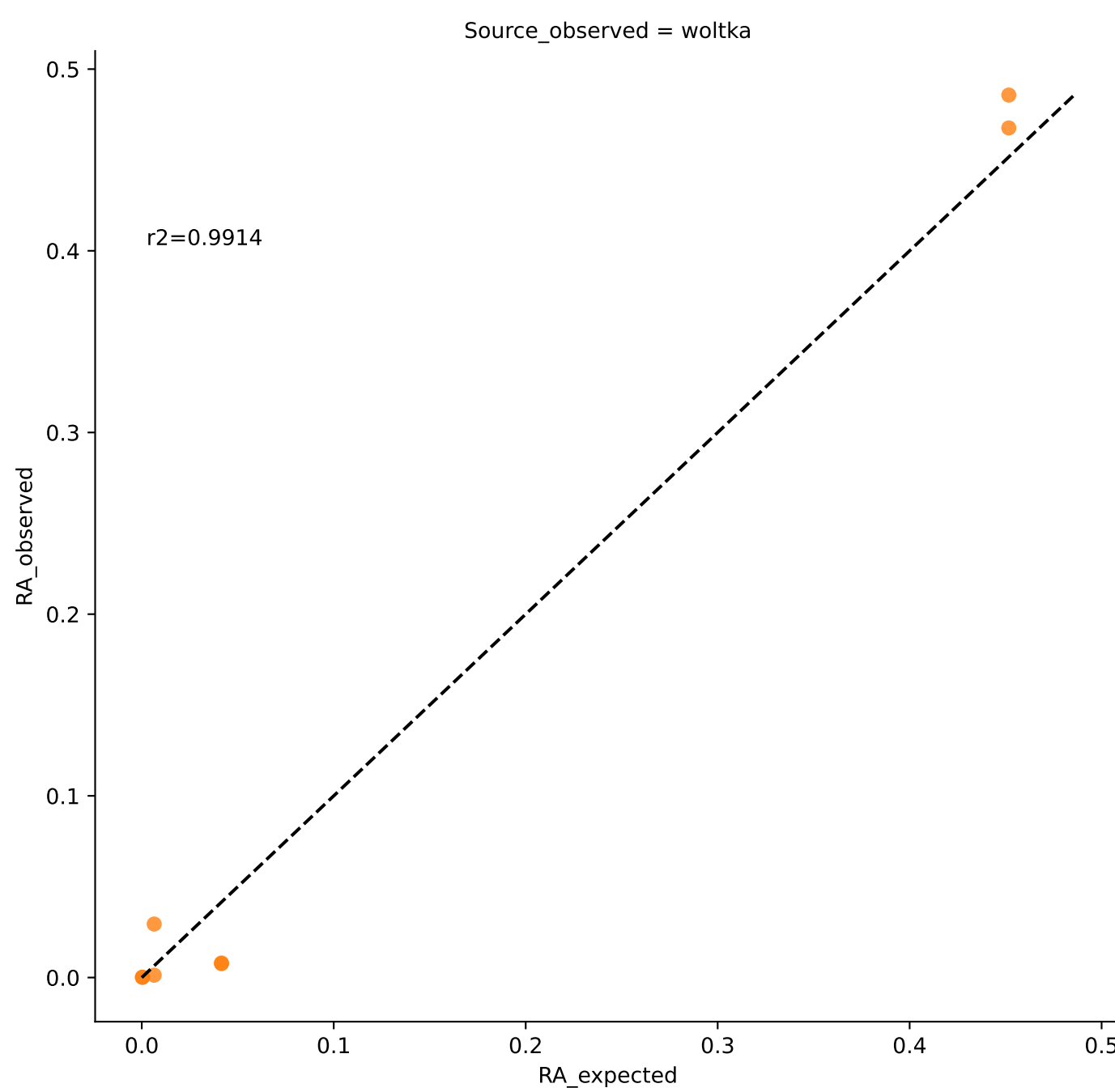
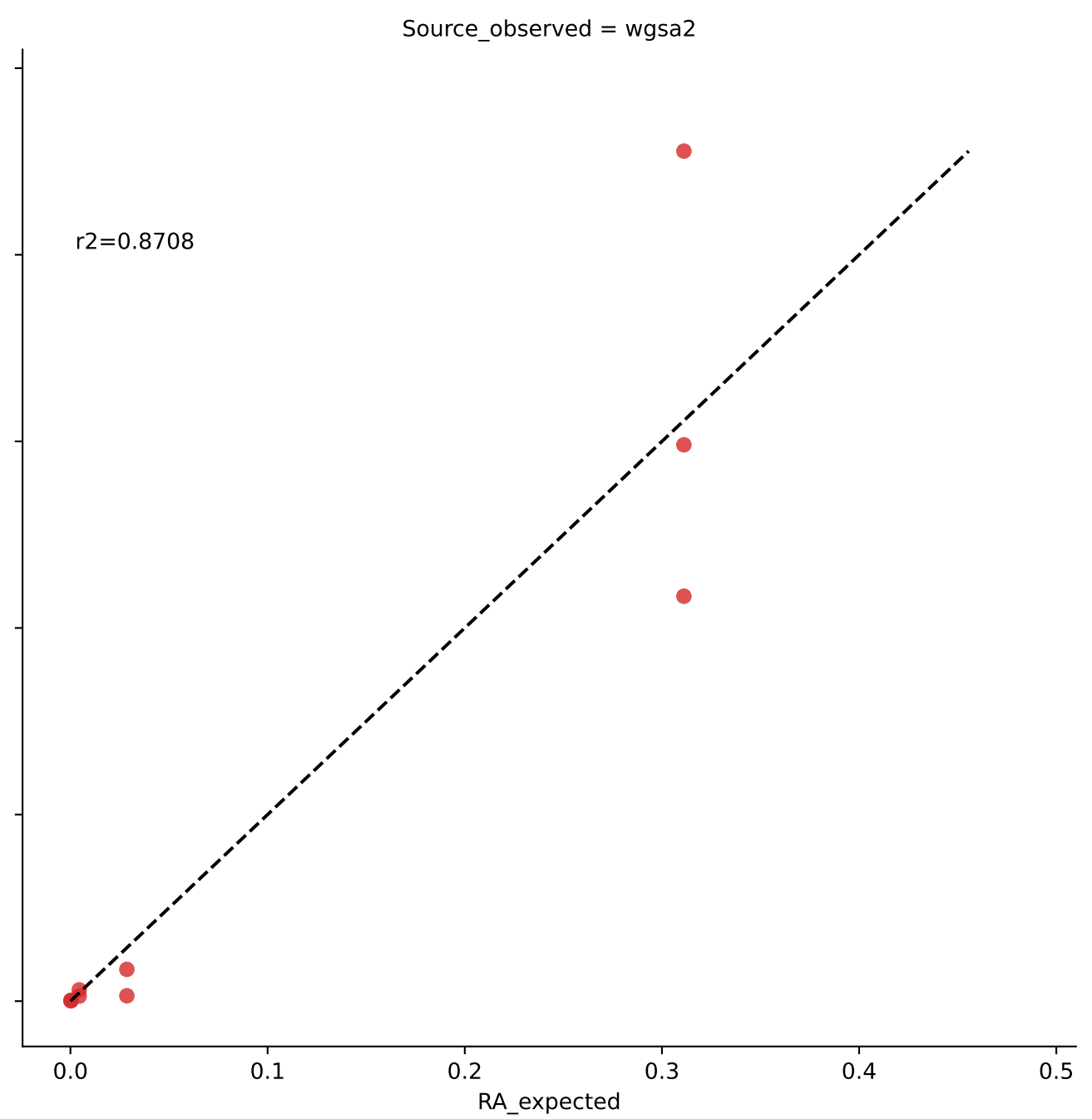
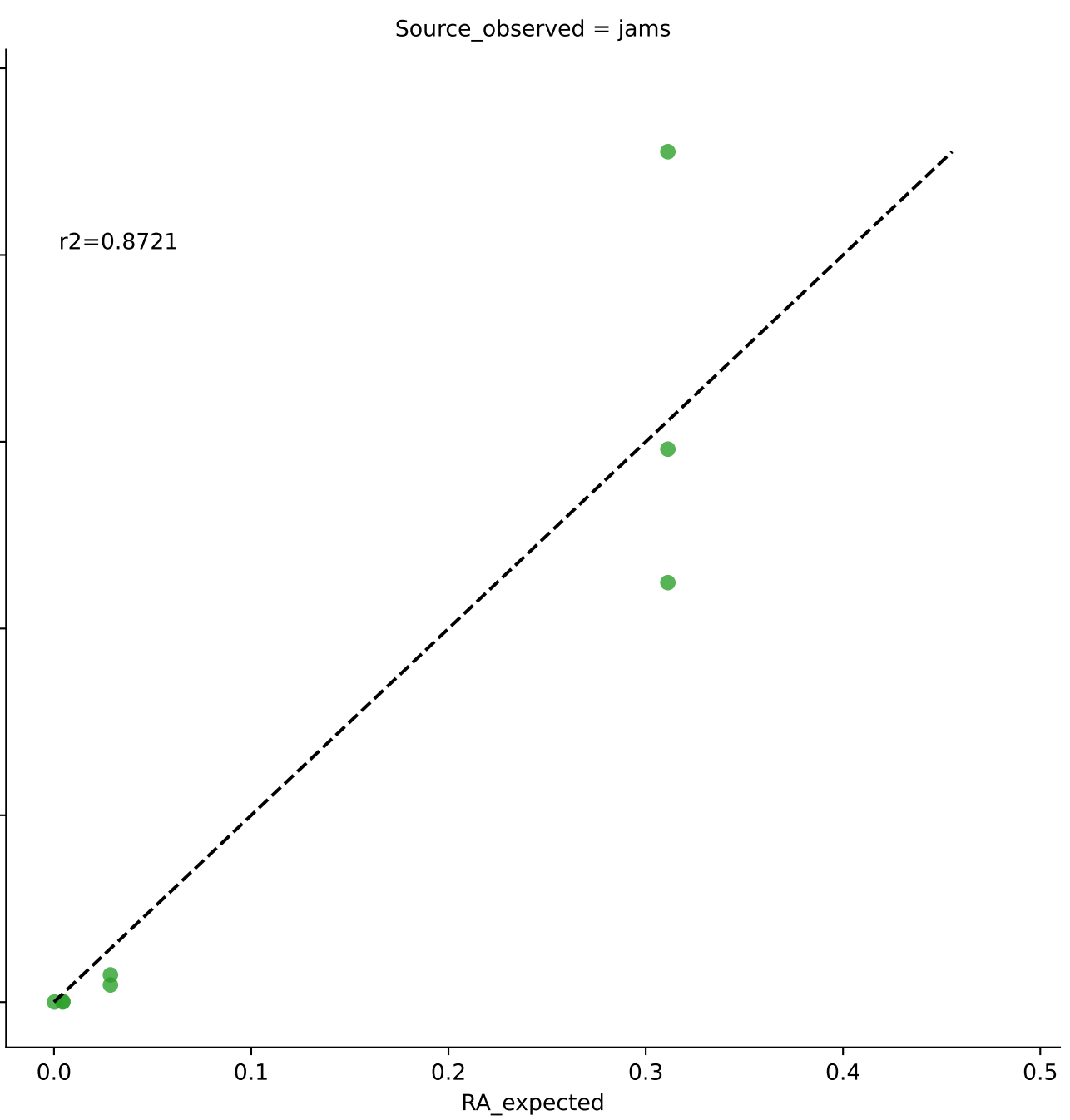
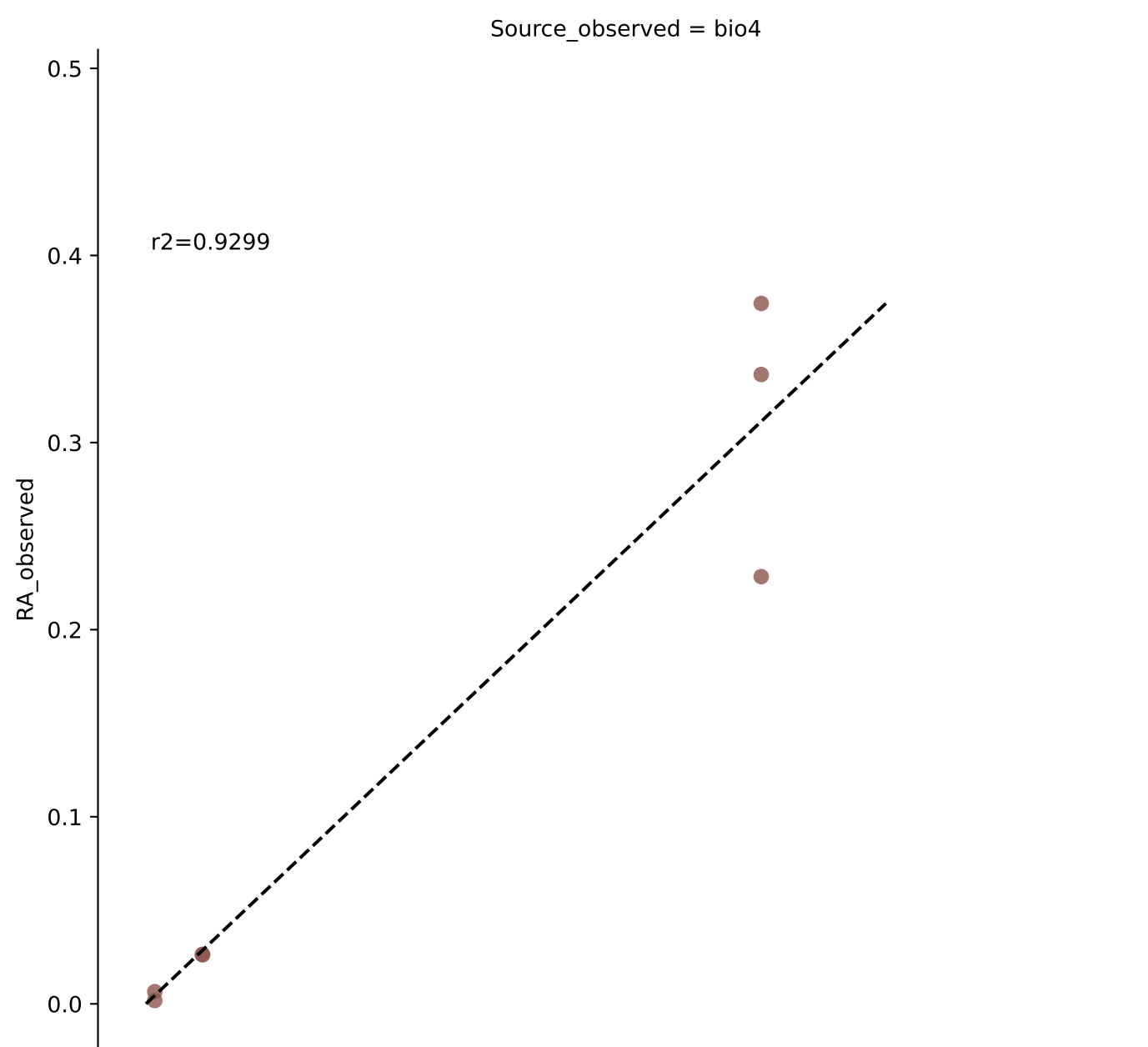


Bivariate Linear Regression for Sample MIX-C in Experiment nist (Species at filter threshold 1e-05)

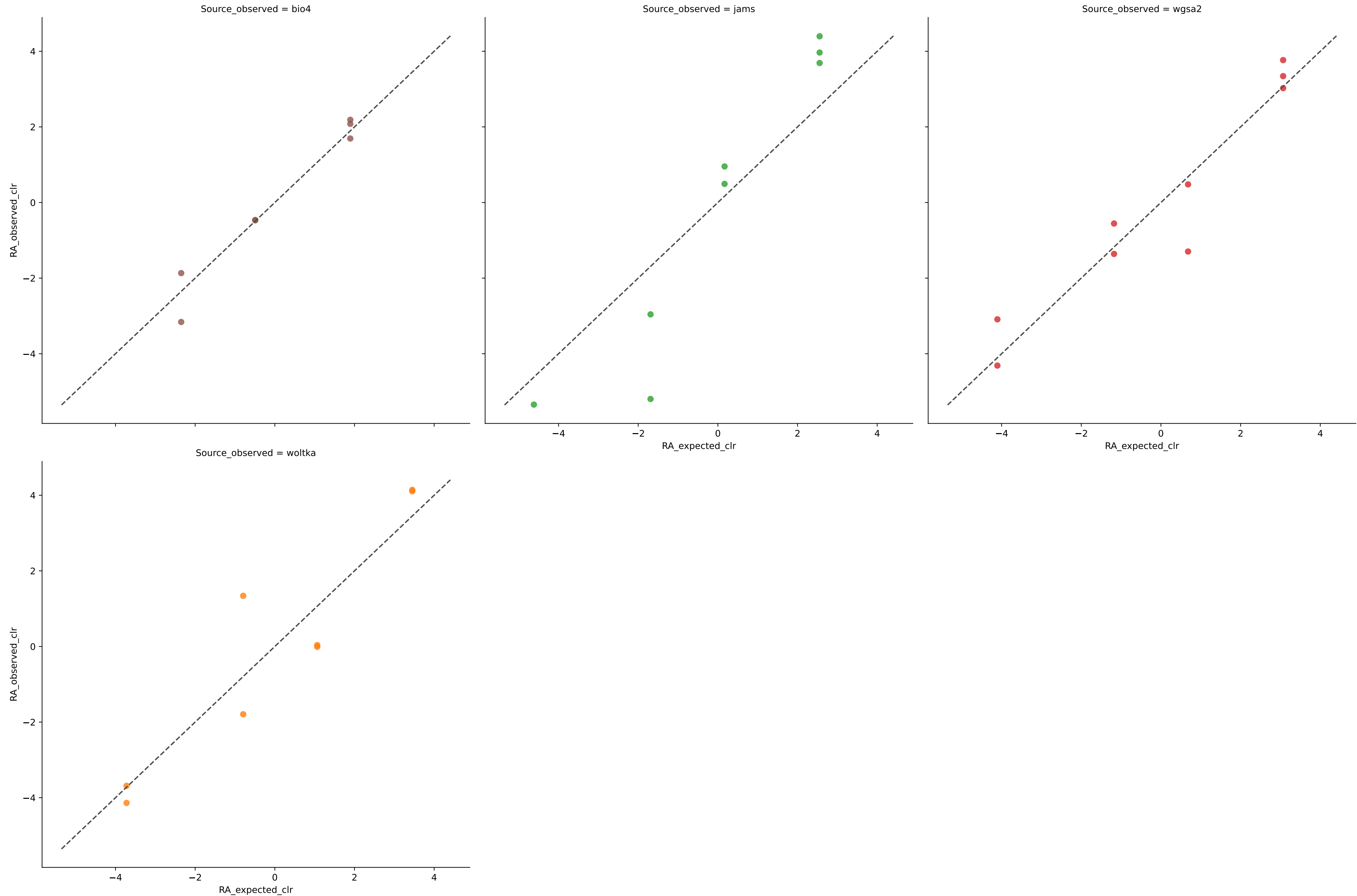


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	7	0.9697	0.0203	0.9148	0.9288	0.0302	100.0000	0.0000
jams	8	0.5064	0.0801	5.6037	0.6794	0.1338	100.0000	0.0000
wgsa2	10	0.3886	0.0791	8.5992	0.6043	0.1476	100.0000	0.0000
woltka	8	0.8059	0.0522	3.0224	0.7911	0.0824	100.0000	0.0000

Bivariate Linear Regression for Sample MIX-D in Experiment nist (Species at filter threshold 1e-05)

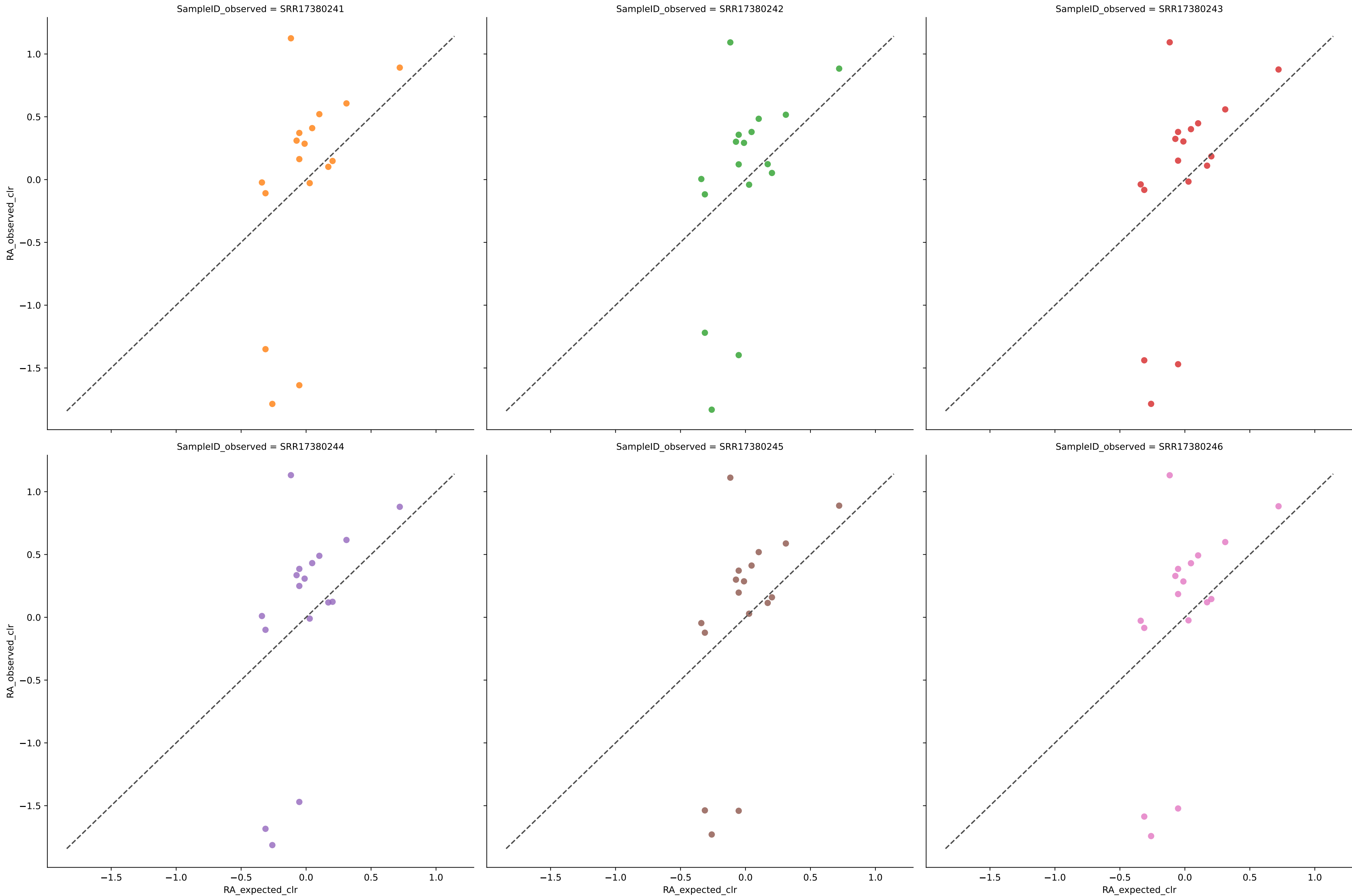


Bivariate Linear Regression for Sample MIX-D in Experiment nist (Species at filter threshold 1e-05)



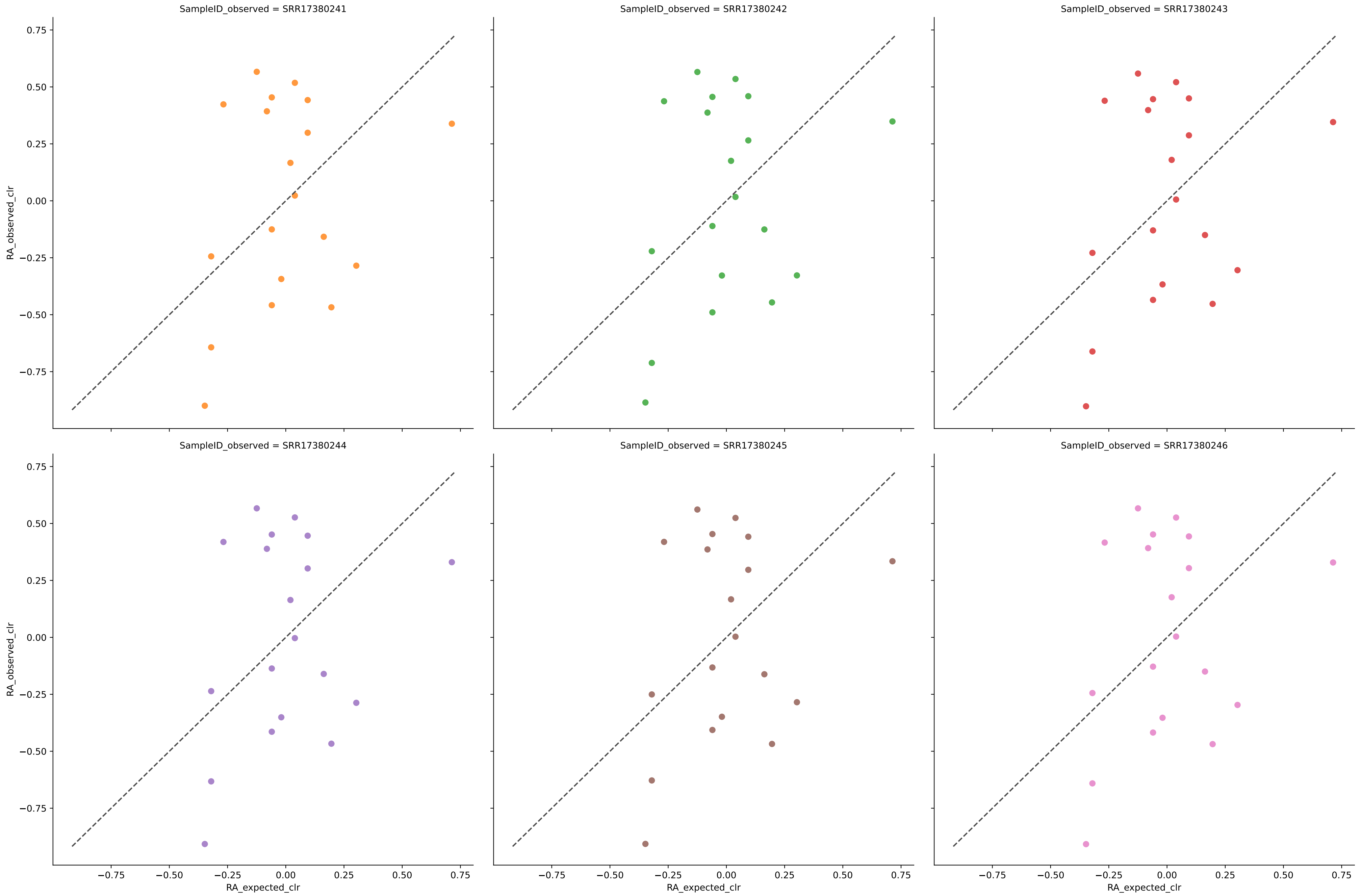
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	7	0.9299	0.0258	1.0255	0.9098	0.0406	100.0000	0.0000
jams	8	0.8721	0.0360	4.6725	0.8559	0.0603	100.0000	0.0000
wgsa2	9	0.8708	0.0325	2.4524	0.8537	0.0584	100.0000	0.0000
woltka	8	0.9914	0.0182	2.9743	0.9270	0.0230	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using bio4 in Experiment tourlousse with filter 1e-05



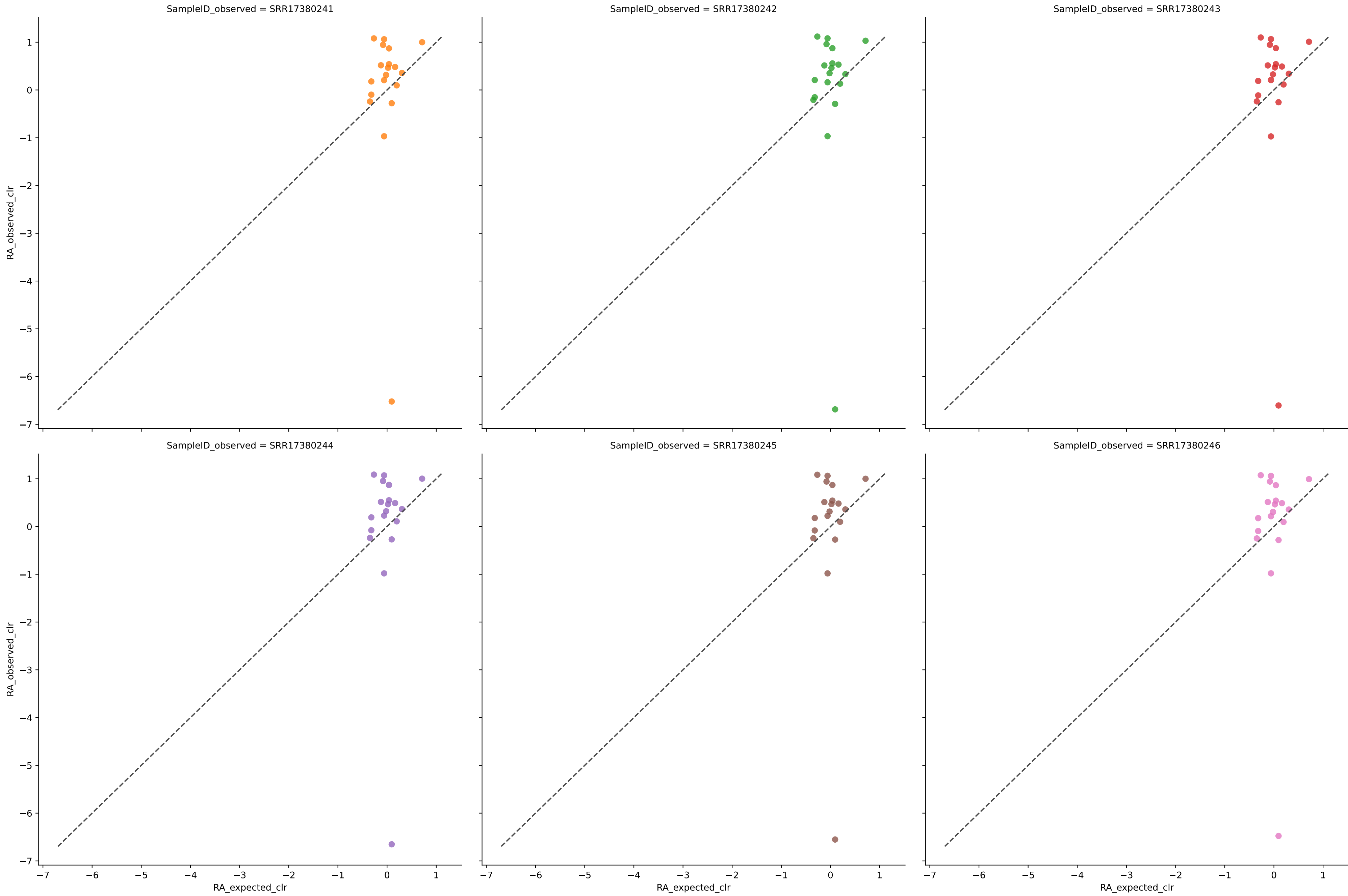
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR17380241	17	0.2683	0.0031	2.9154	0.8404	0.0048	100.0000	0.0000
SRR17380242	17	0.2667	0.0030	2.7401	0.8446	0.0047	100.0000	0.0000
SRR17380243	17	0.2744	0.0030	2.8350	0.8461	0.0046	100.0000	0.0000
SRR17380244	17	0.2551	0.0032	3.0019	0.8349	0.0048	100.0000	0.0000
SRR17380245	17	0.2771	0.0031	2.8973	0.8429	0.0047	100.0000	0.0000
SRR17380246	17	0.2608	0.0032	2.9331	0.8386	0.0048	100.0000	0.0000
Average	17	0.2671	0.0031	2.8871	0.8412	0.0048	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using jams in Experiment tourlousse with filter 1e-05



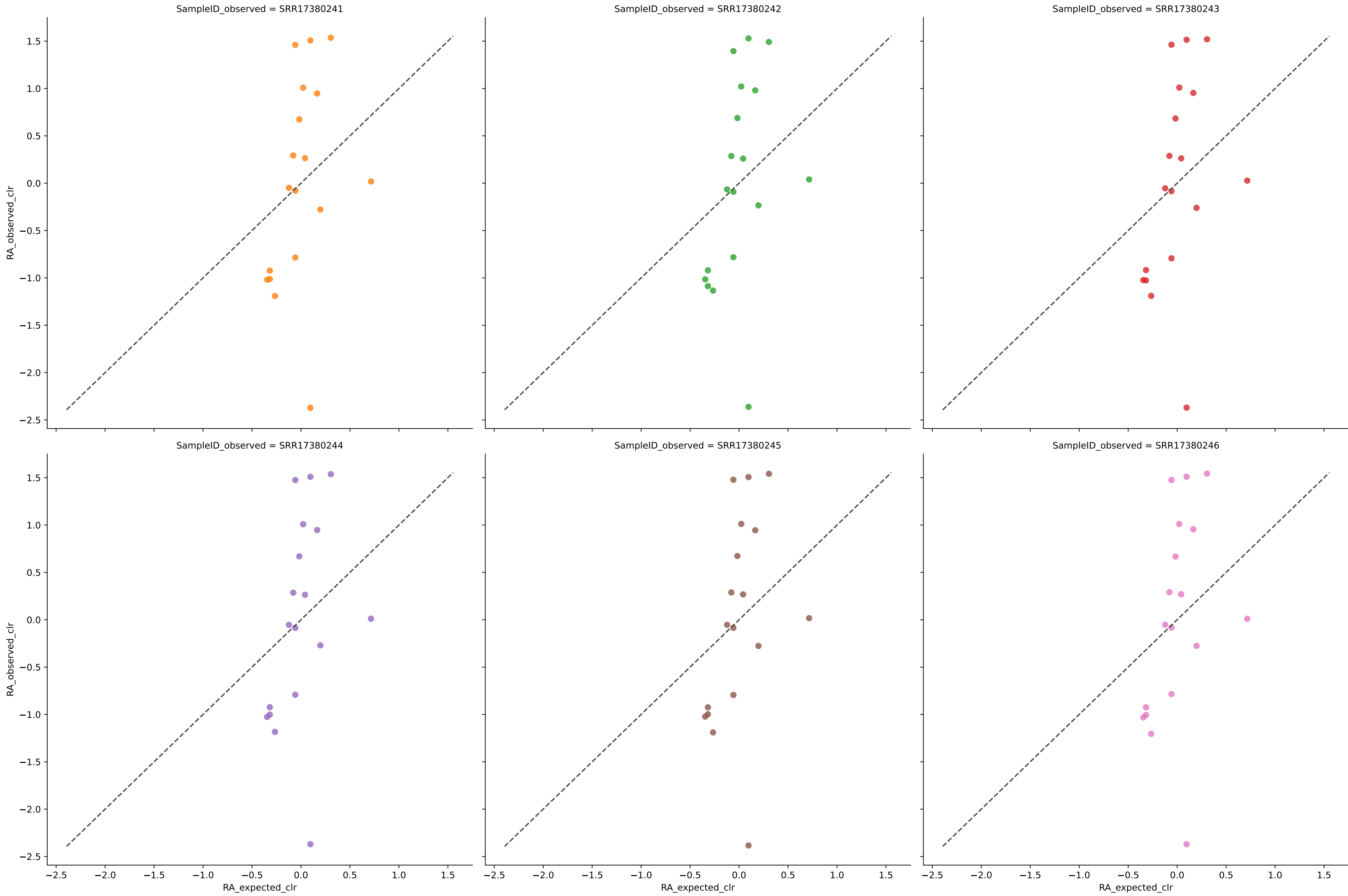
	Diversity	R ²	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR17380241	19	0.0294	0.0033	1.8893	0.8104	0.0039	100.0000	0.0000
SRR17380242	19	0.0298	0.0033	1.9110	0.8093	0.0039	100.0000	0.0000
SRR17380243	19	0.0296	0.0033	1.8964	0.8091	0.0039	100.0000	0.0000
SRR17380244	19	0.0276	0.0033	1.8859	0.8093	0.0039	100.0000	0.0000
SRR17380245	19	0.0291	0.0033	1.8776	0.8108	0.0039	100.0000	0.0000
SRR17380246	19	0.0276	0.0033	1.8902	0.8094	0.0039	100.0000	0.0000
Average	19	0.0289	0.0033	1.8917	0.8097	0.0039	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using wgsa in Experiment tourlousse with filter 1e-05



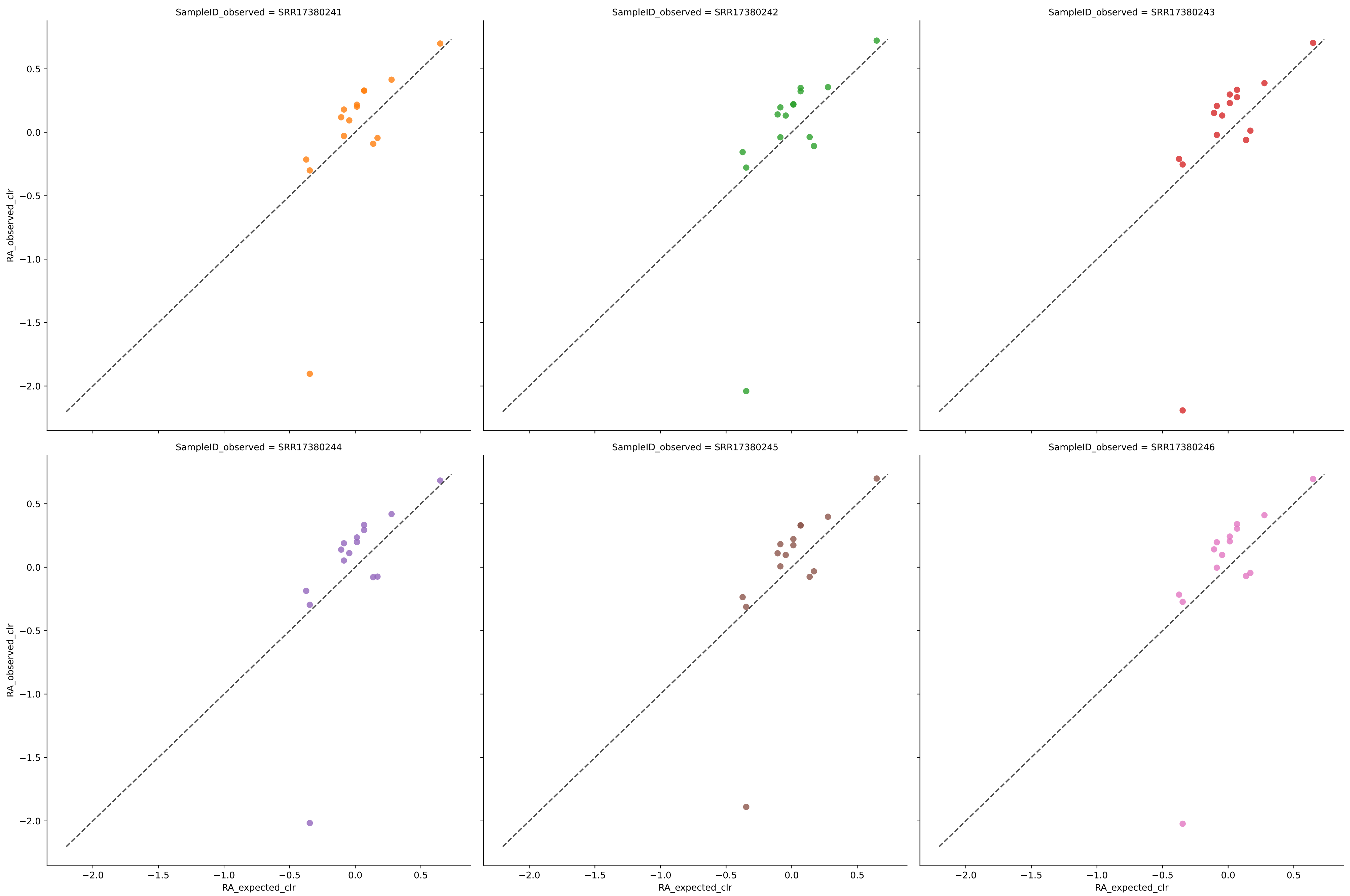
	Diversity	R ²	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR17380241	19	0.0476	0.0036	7.1479	0.7940	0.0048	100.0000	0.0000
SRR17380242	19	0.0495	0.0036	7.3192	0.7927	0.0048	100.0000	0.0000
SRR17380243	19	0.0477	0.0036	7.2308	0.7938	0.0048	100.0000	0.0000
SRR17380244	19	0.0464	0.0036	7.2784	0.7946	0.0048	100.0000	0.0000
SRR17380245	19	0.0474	0.0036	7.1787	0.7948	0.0048	100.0000	0.0000
SRR17380246	19	0.0471	0.0036	7.1059	0.7944	0.0048	100.0000	0.0000
Average	19	0.0476	0.0036	7.2102	0.7941	0.0048	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using woltka in Experiment tourlousse with filter 1e-05



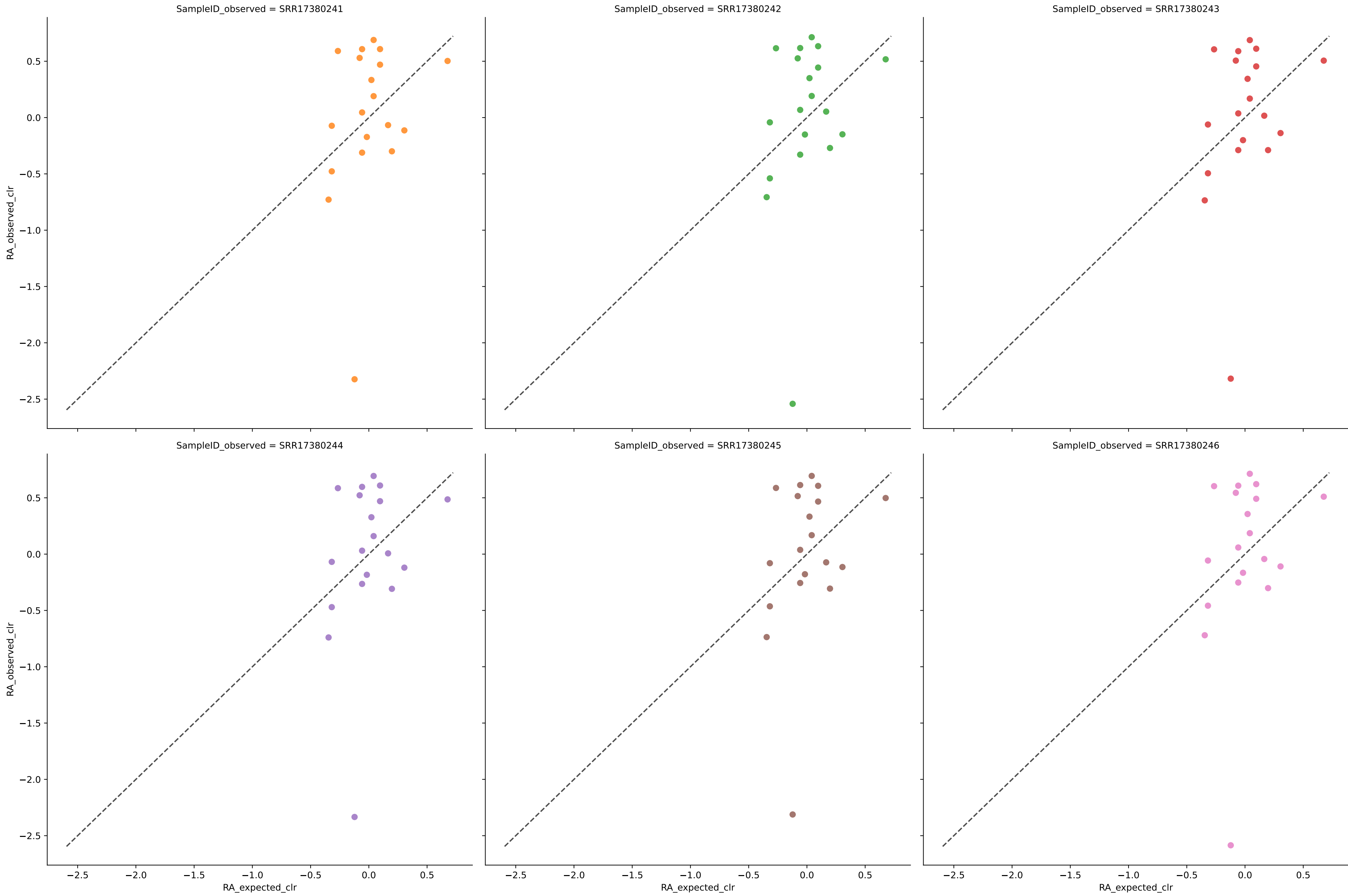
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR17380241	18	0.0619	0.0067	4.1905	0.6387	0.0082	100.0000	0.0000
SRR17380242	18	0.0655	0.0066	4.1572	0.6414	0.0081	100.0000	0.0000
SRR17380243	18	0.0618	0.0067	4.1904	0.6385	0.0082	100.0000	0.0000
SRR17380244	18	0.0608	0.0067	4.1939	0.6374	0.0083	100.0000	0.0000
SRR17380245	18	0.0609	0.0067	4.2048	0.6373	0.0083	100.0000	0.0000
SRR17380246	18	0.0611	0.0067	4.2050	0.6370	0.0083	100.0000	0.0000
Average	18	0.0620	0.0067	4.1903	0.6384	0.0082	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using bio4 in Experiment tourlousse with filter 1e-05



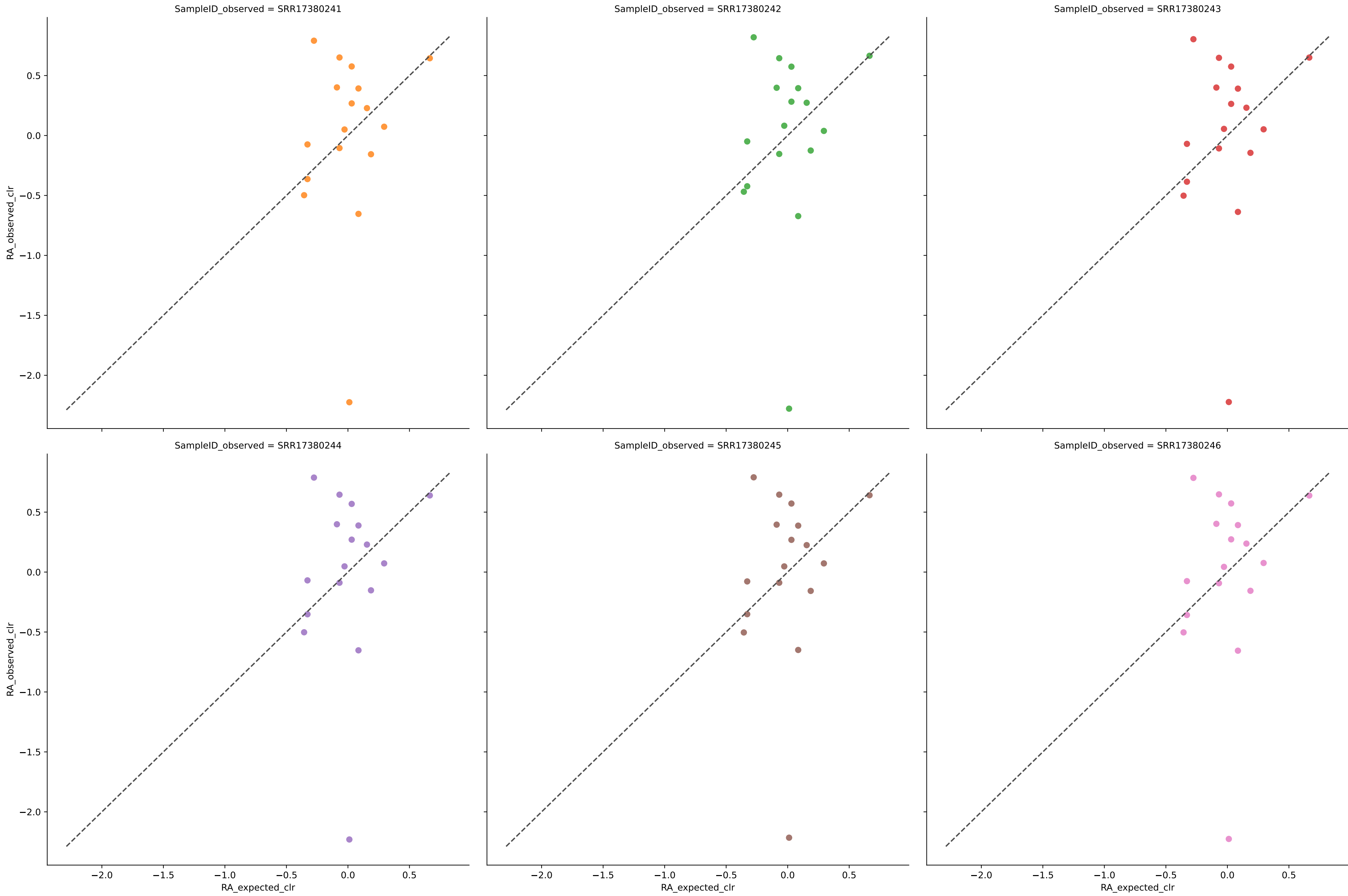
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR17380241	15	0.6588	0.0018	1.7116	0.9196	0.0023	100.0000	0.0000
SRR17380242	15	0.6349	0.0018	1.8567	0.9185	0.0024	100.0000	0.0000
SRR17380243	15	0.6416	0.0018	1.9887	0.9203	0.0024	100.0000	0.0000
SRR17380244	15	0.6284	0.0019	1.8273	0.9153	0.0024	100.0000	0.0000
SRR17380245	15	0.6673	0.0017	1.6908	0.9221	0.0023	100.0000	0.0000
SRR17380246	15	0.6450	0.0018	1.8257	0.9199	0.0024	100.0000	0.0000
Average	15	0.6460	0.0018	1.8168	0.9193	0.0024	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using jams in Experiment tourlousse with filter 1e-05



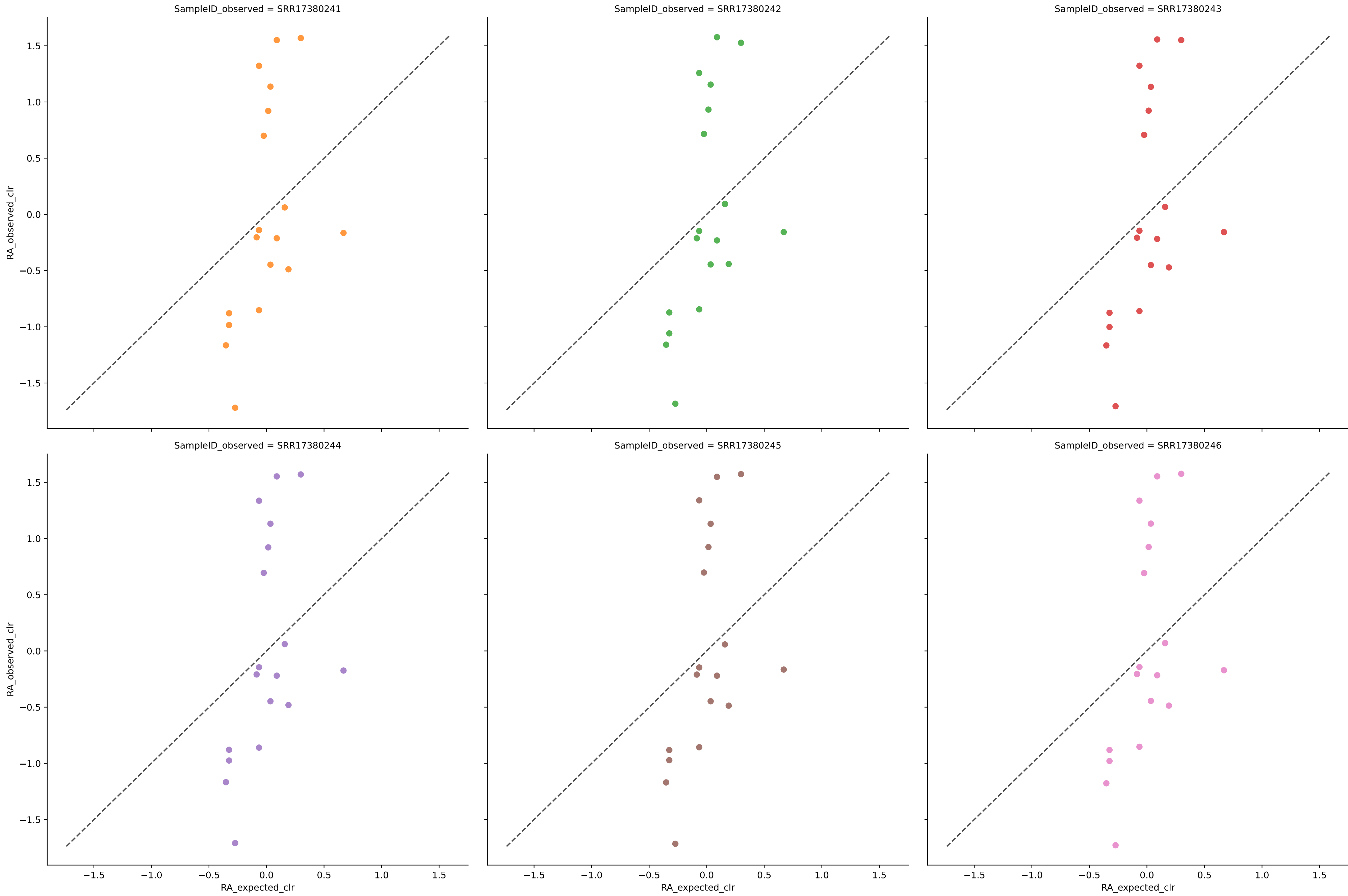
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR17380241	19	0.0685	0.0035	2.8585	0.8028	0.0040	100.0000	0.0000
SRR17380242	19	0.0699	0.0034	3.0443	0.8034	0.0041	100.0000	0.0000
SRR17380243	19	0.0715	0.0034	2.8464	0.8049	0.0040	100.0000	0.0000
SRR17380244	19	0.0671	0.0034	2.8565	0.8044	0.0040	100.0000	0.0000
SRR17380245	19	0.0674	0.0034	2.8435	0.8038	0.0040	100.0000	0.0000
SRR17380246	19	0.0661	0.0035	3.0762	0.8022	0.0041	100.0000	0.0000
Average	19	0.0684	0.0034	2.9209	0.8036	0.0040	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using wgsa in Experiment tourlousse with filter 1e-05



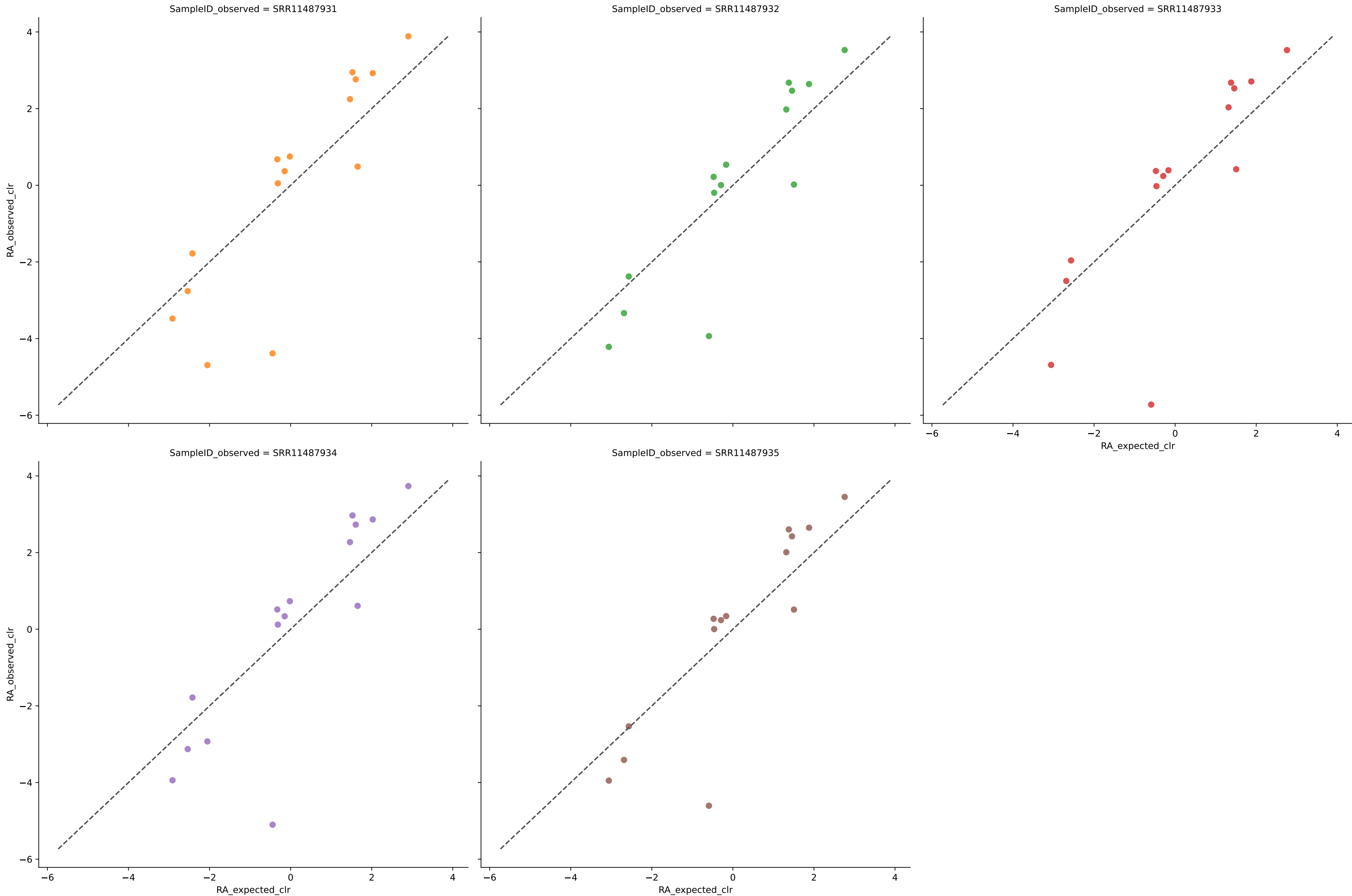
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR17380241	17	0.0588	0.0036	2.8562	0.8157	0.0047	100.0000	0.0000
SRR17380242	17	0.0603	0.0036	2.9180	0.8144	0.0047	100.0000	0.0000
SRR17380243	17	0.0586	0.0036	2.8552	0.8154	0.0047	100.0000	0.0000
SRR17380244	17	0.0579	0.0036	2.8562	0.8166	0.0047	100.0000	0.0000
SRR17380245	17	0.0583	0.0036	2.8440	0.8167	0.0047	100.0000	0.0000
SRR17380246	17	0.0586	0.0036	2.8549	0.8162	0.0047	100.0000	0.0000
Average	17	0.0587	0.0036	2.8641	0.8158	0.0047	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using woltka in Experiment tourlousse with filter 1e-05



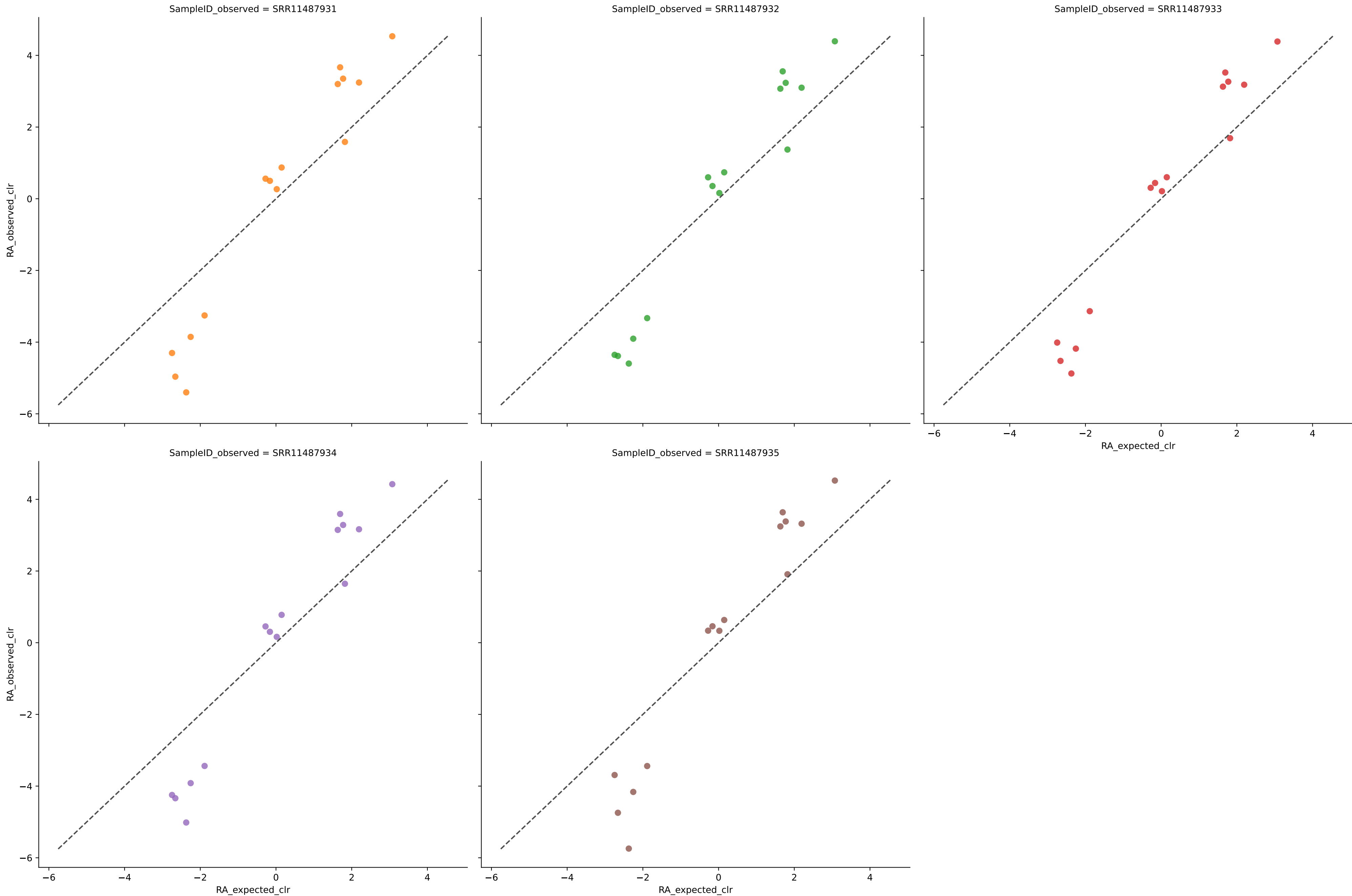
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR17380241	18	0.0560	0.0073	3.7226	0.6054	0.0086	100.0000	0.0000
SRR17380242	18	0.0568	0.0072	3.6942	0.6082	0.0084	100.0000	0.0000
SRR17380243	18	0.0555	0.0073	3.7162	0.6058	0.0086	100.0000	0.0000
SRR17380244	18	0.0550	0.0073	3.7255	0.6044	0.0086	100.0000	0.0000
SRR17380245	18	0.0554	0.0073	3.7281	0.6042	0.0086	100.0000	0.0000
SRR17380246	18	0.0555	0.0073	3.7368	0.6044	0.0086	100.0000	0.0000
Average	18	0.0557	0.0073	3.7206	0.6054	0.0086	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using bio4 in Experiment Amos hilo with filter 1e-05



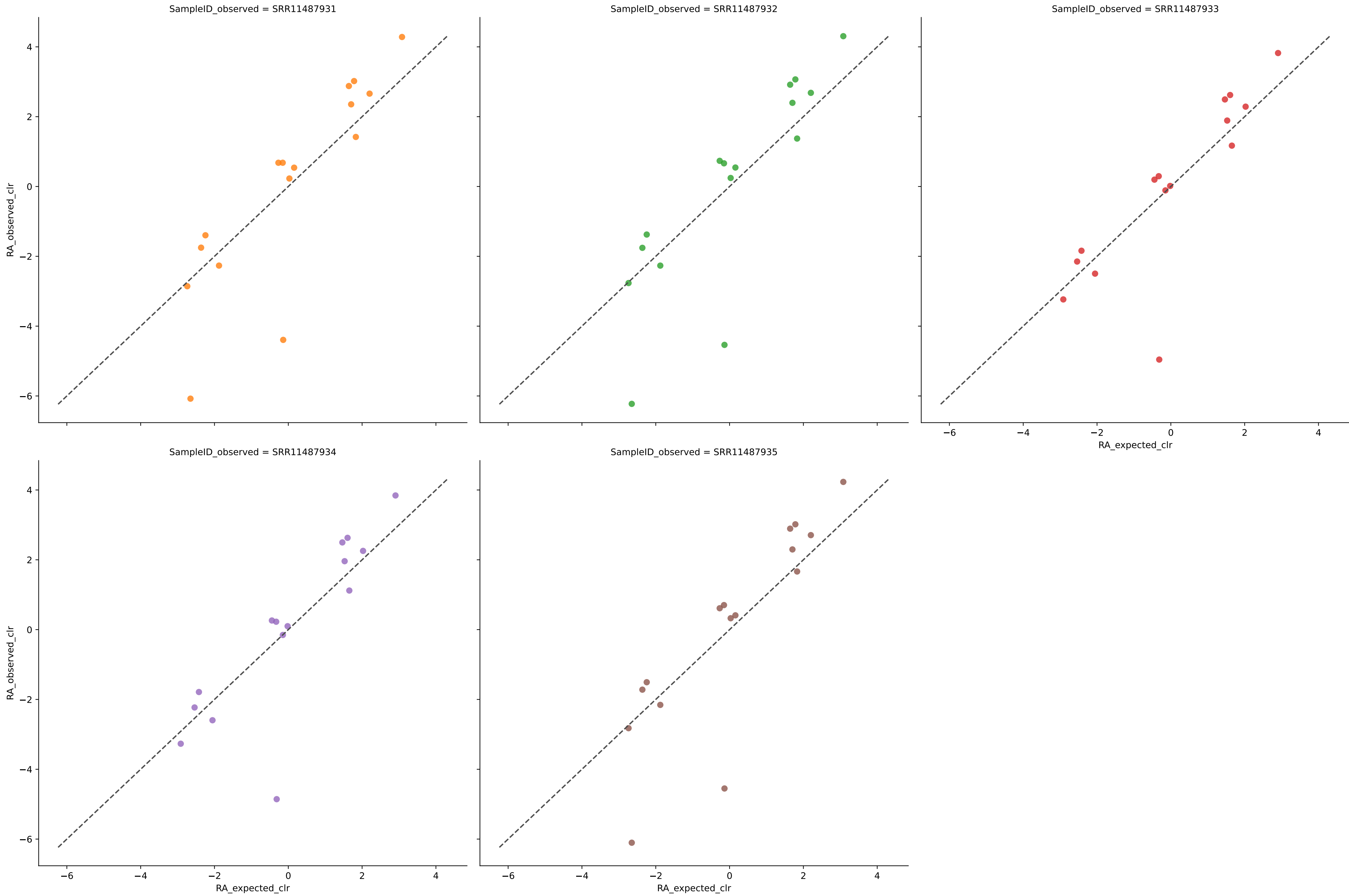
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487931	15	0.9163	0.0035	5.6916	0.8671	0.0064	100.0000	0.0000
SRR11487932	14	0.9020	0.0037	4.5410	0.8713	0.0067	100.0000	0.0000
SRR11487933	14	0.9056	0.0034	6.0565	0.8801	0.0064	100.0000	0.0000
SRR11487934	15	0.8969	0.0032	5.6939	0.8806	0.0065	100.0000	0.0000
SRR11487935	14	0.9105	0.0032	4.8690	0.8881	0.0061	100.0000	0.0000
Average	14	0.9063	0.0034	5.3704	0.8774	0.0064	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using jams in Experiment Amos hilo with filter 1e-05



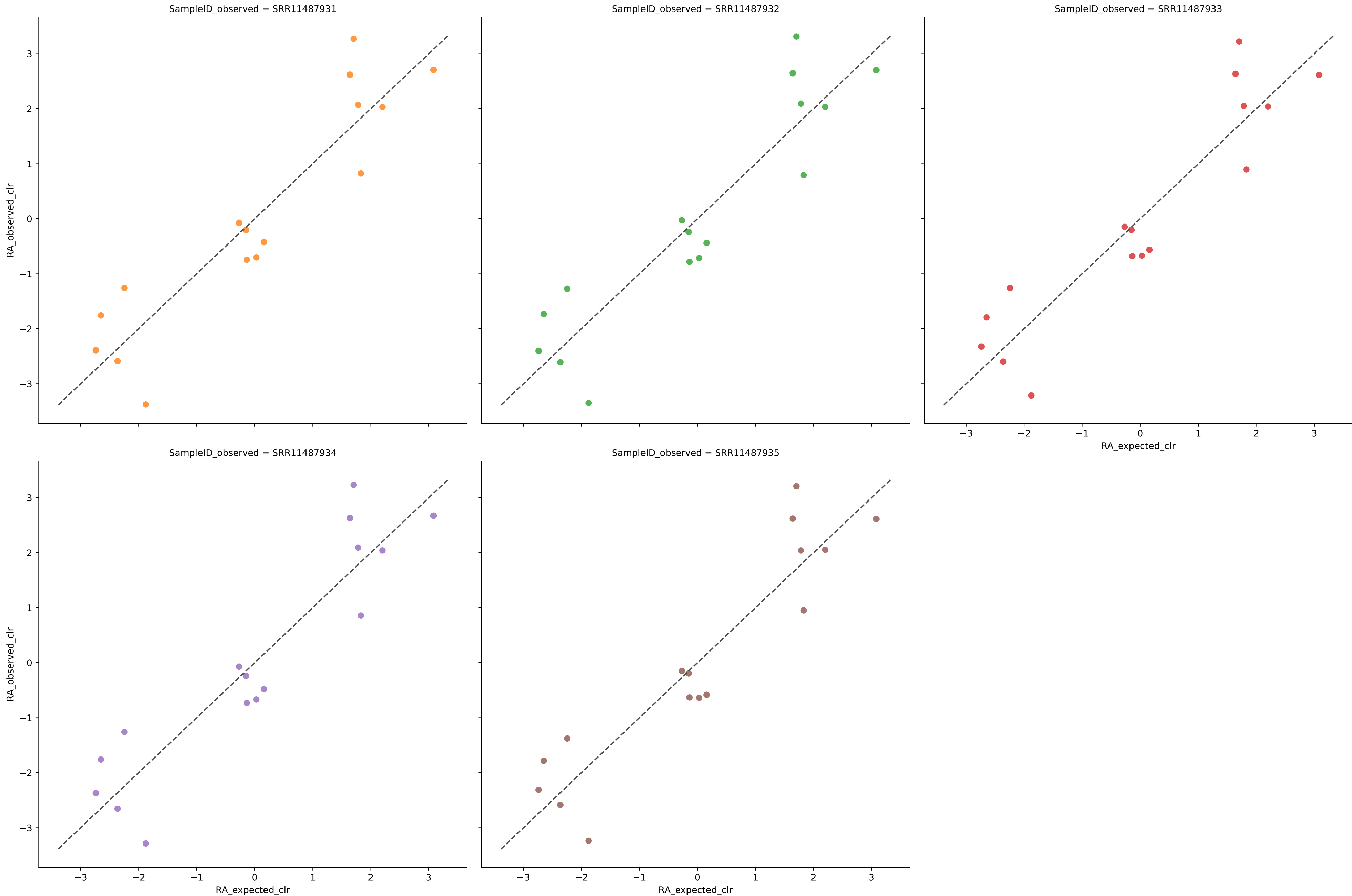
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487931	15	0.9019	0.0045	5.9316	0.8311	0.0071	100.0000	0.0000
SRR11487932	15	0.8955	0.0046	5.2105	0.8292	0.0072	100.0000	0.0000
SRR11487933	15	0.9096	0.0042	5.2991	0.8409	0.0065	100.0000	0.0000
SRR11487934	15	0.9006	0.0044	5.4209	0.8340	0.0069	100.0000	0.0000
SRR11487935	15	0.9155	0.0042	6.0090	0.8426	0.0064	100.0000	0.0000
Average	15	0.9047	0.0044	5.5742	0.8356	0.0068	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using wgsa in Experiment Amos hilo with filter 1e-05



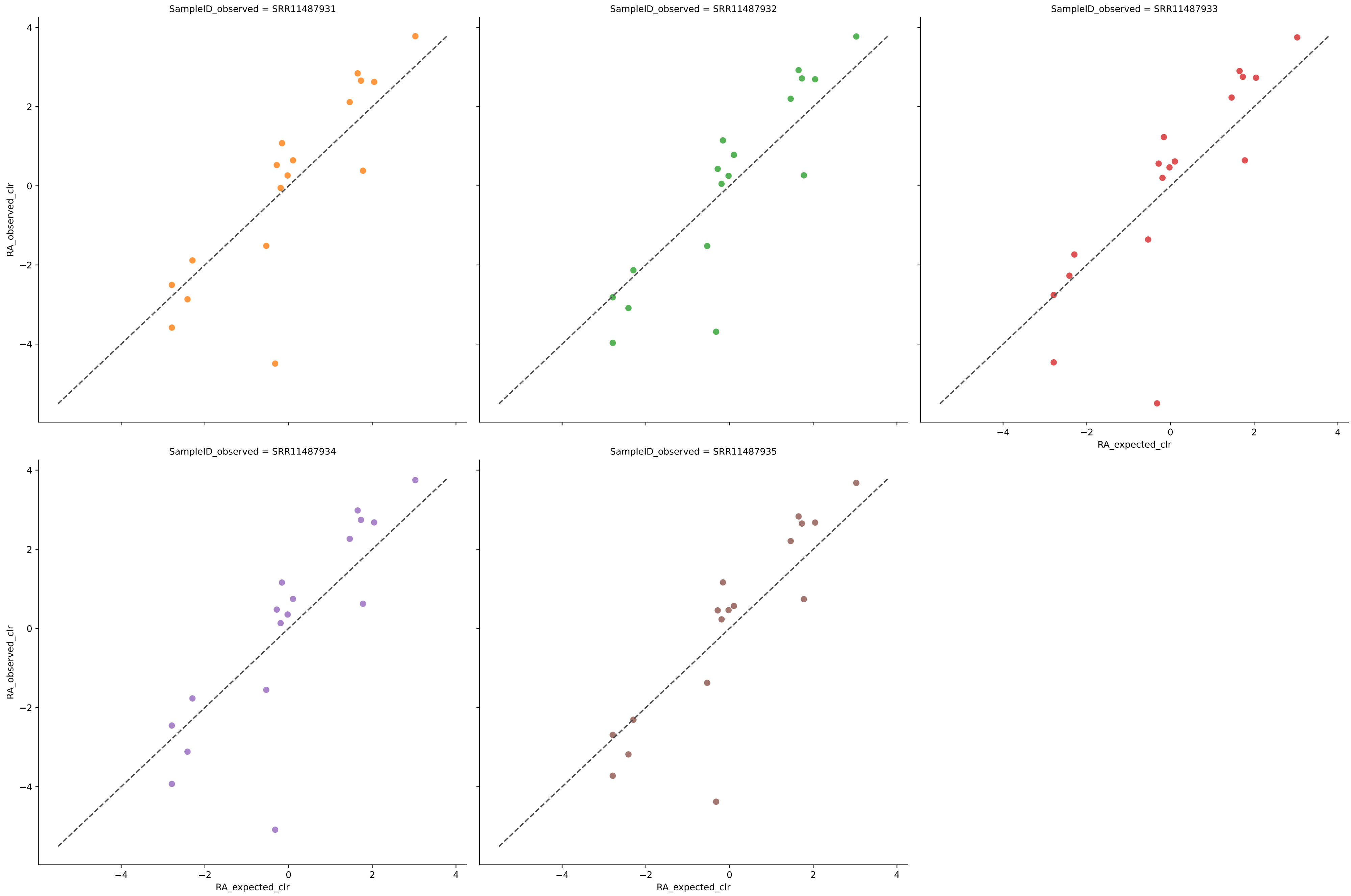
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487931	16	0.9134	0.0048	6.1797	0.8064	0.0083	100.0000	0.0000
SRR11487932	16	0.9112	0.0049	6.4008	0.8047	0.0082	100.0000	0.0000
SRR11487933	15	0.9190	0.0049	5.1489	0.8147	0.0079	100.0000	0.0000
SRR11487934	15	0.9156	0.0050	5.0880	0.8122	0.0081	100.0000	0.0000
SRR11487935	16	0.9235	0.0046	6.2562	0.8180	0.0075	100.0000	0.0000
Average	16	0.9165	0.0048	5.8147	0.8112	0.0080	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using wol in Experiment Amos hilo with filter 1e-05



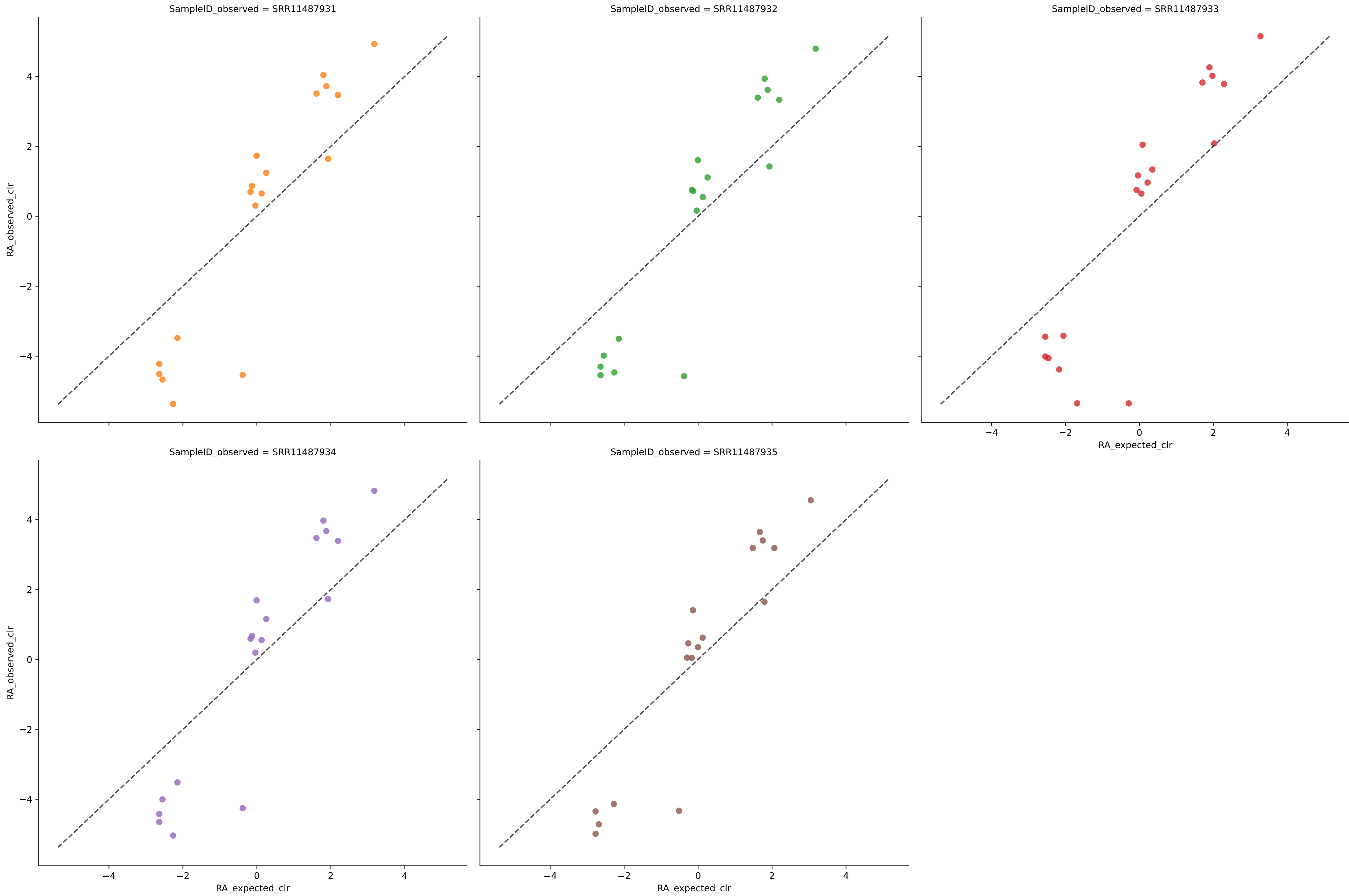
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487931	16	0.3723	0.0086	3.1882	0.6543	0.0166	100.0000	0.0000
SRR11487932	16	0.3532	0.0088	3.2368	0.6454	0.0171	100.0000	0.0000
SRR11487933	16	0.3586	0.0087	3.0849	0.6509	0.0168	100.0000	0.0000
SRR11487934	16	0.3760	0.0086	3.1331	0.6548	0.0165	100.0000	0.0000
SRR11487935	16	0.3660	0.0086	3.0234	0.6564	0.0166	100.0000	0.0000
Average	16	0.3652	0.0087	3.1333	0.6524	0.0167	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using bio4 in Experiment Amos hilo with filter 1e-05



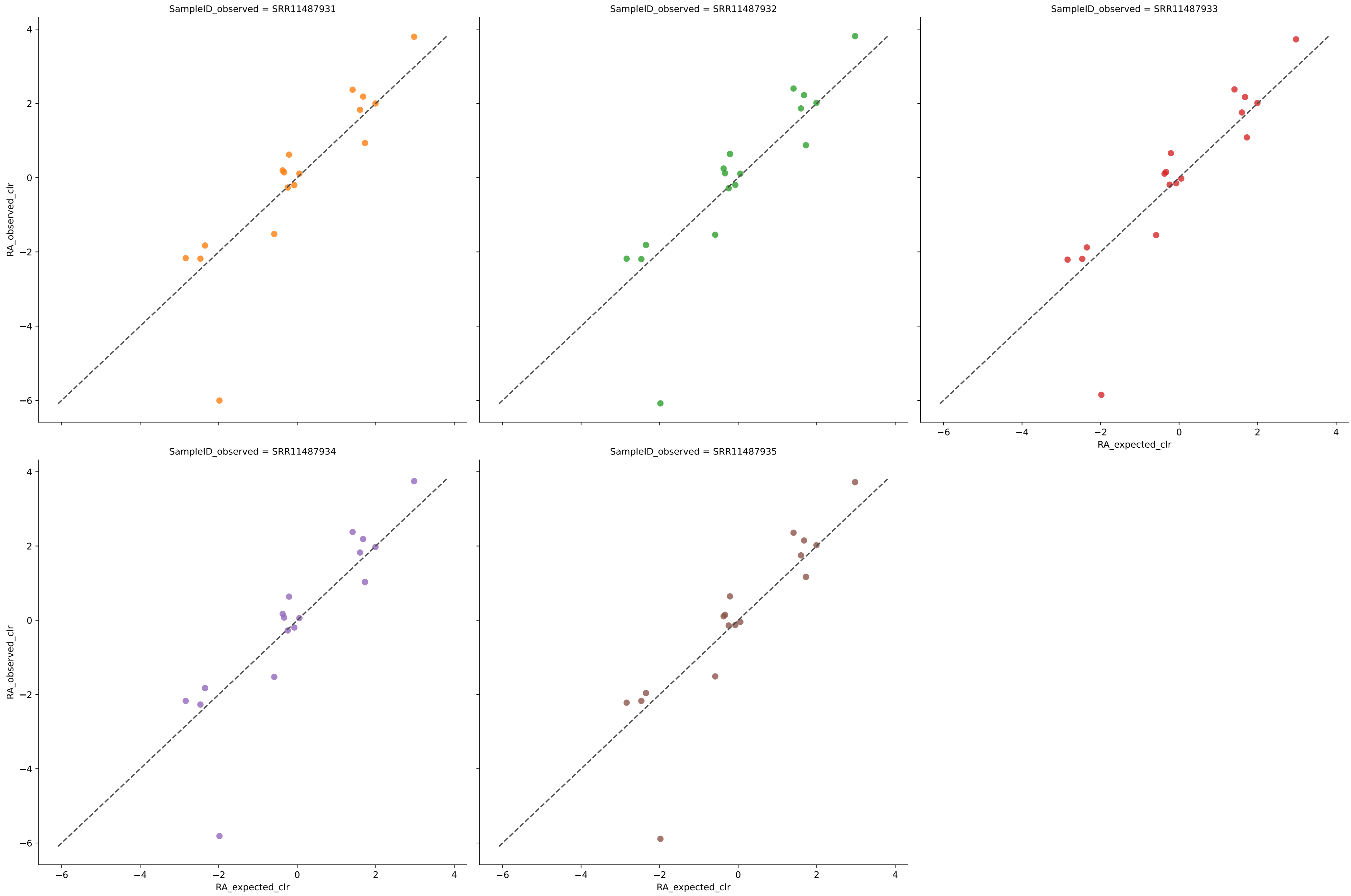
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487931	17	0.9142	0.0034	5.2528	0.8570	0.0060	100.0000	0.0000
SRR11487932	17	0.9028	0.0033	4.8354	0.8613	0.0062	100.0000	0.0000
SRR11487933	17	0.9066	0.0030	6.2762	0.8717	0.0058	100.0000	0.0000
SRR11487934	17	0.8944	0.0032	5.8814	0.8635	0.0062	100.0000	0.0000
SRR11487935	17	0.9113	0.0029	5.1244	0.8775	0.0056	100.0000	0.0000
Average	17	0.9059	0.0031	5.4740	0.8662	0.0060	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using jams in Experiment Amos hilo with filter 1e-05



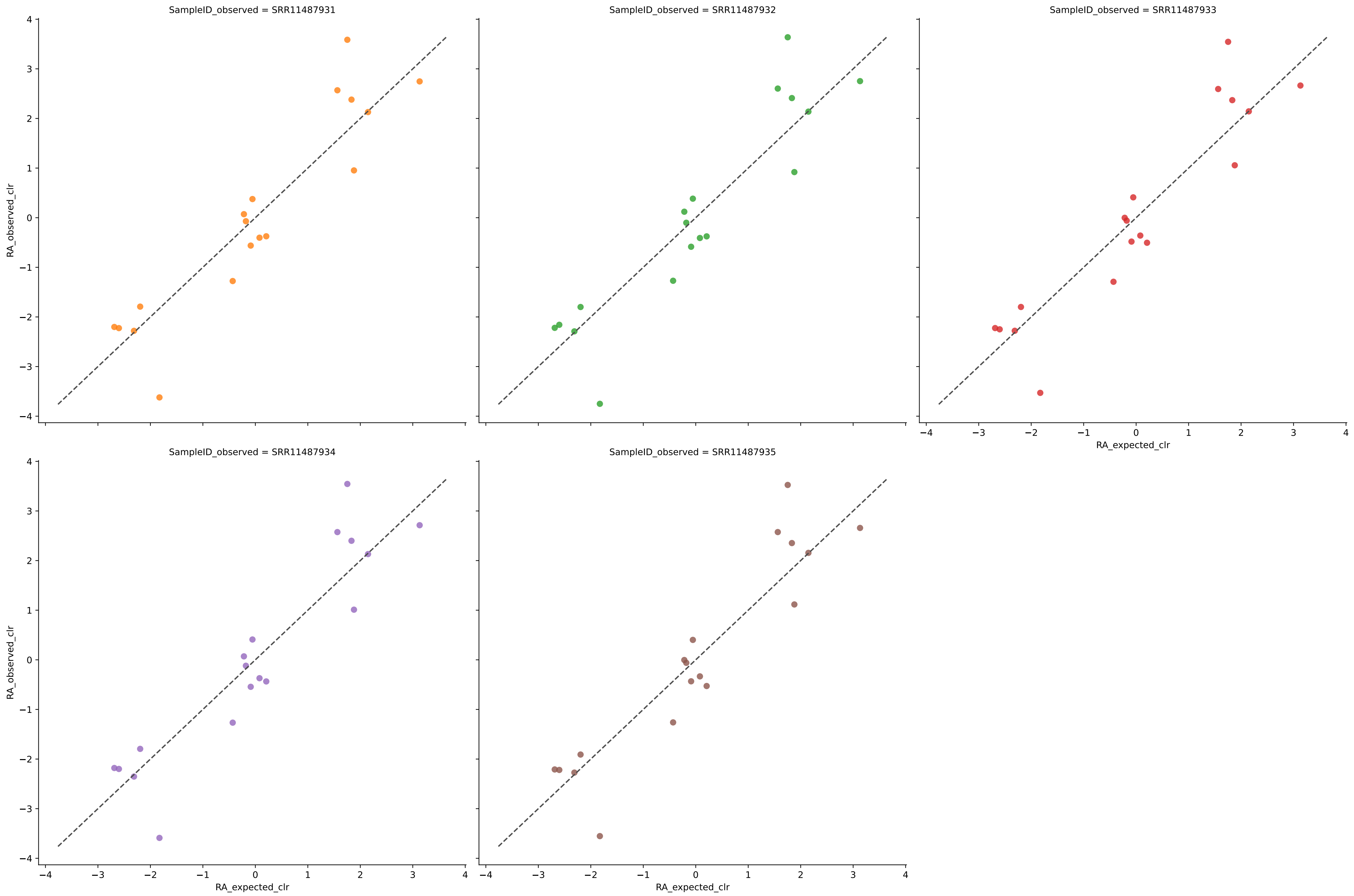
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487931	18	0.9007	0.0043	7.8716	0.8087	0.0069	100.0000	0.0000
SRR11487932	18	0.8945	0.0043	7.2568	0.8062	0.0070	100.0000	0.0000
SRR11487933	19	0.9069	0.0039	8.8944	0.8148	0.0063	100.0000	0.0000
SRR11487934	18	0.8980	0.0042	7.3804	0.8101	0.0067	100.0000	0.0000
SRR11487935	17	0.9106	0.0042	6.7833	0.8194	0.0065	100.0000	0.0000
Average	18	0.9021	0.0042	7.6373	0.8118	0.0067	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using wgsa in Experiment Amos hilo with filter 1e-05



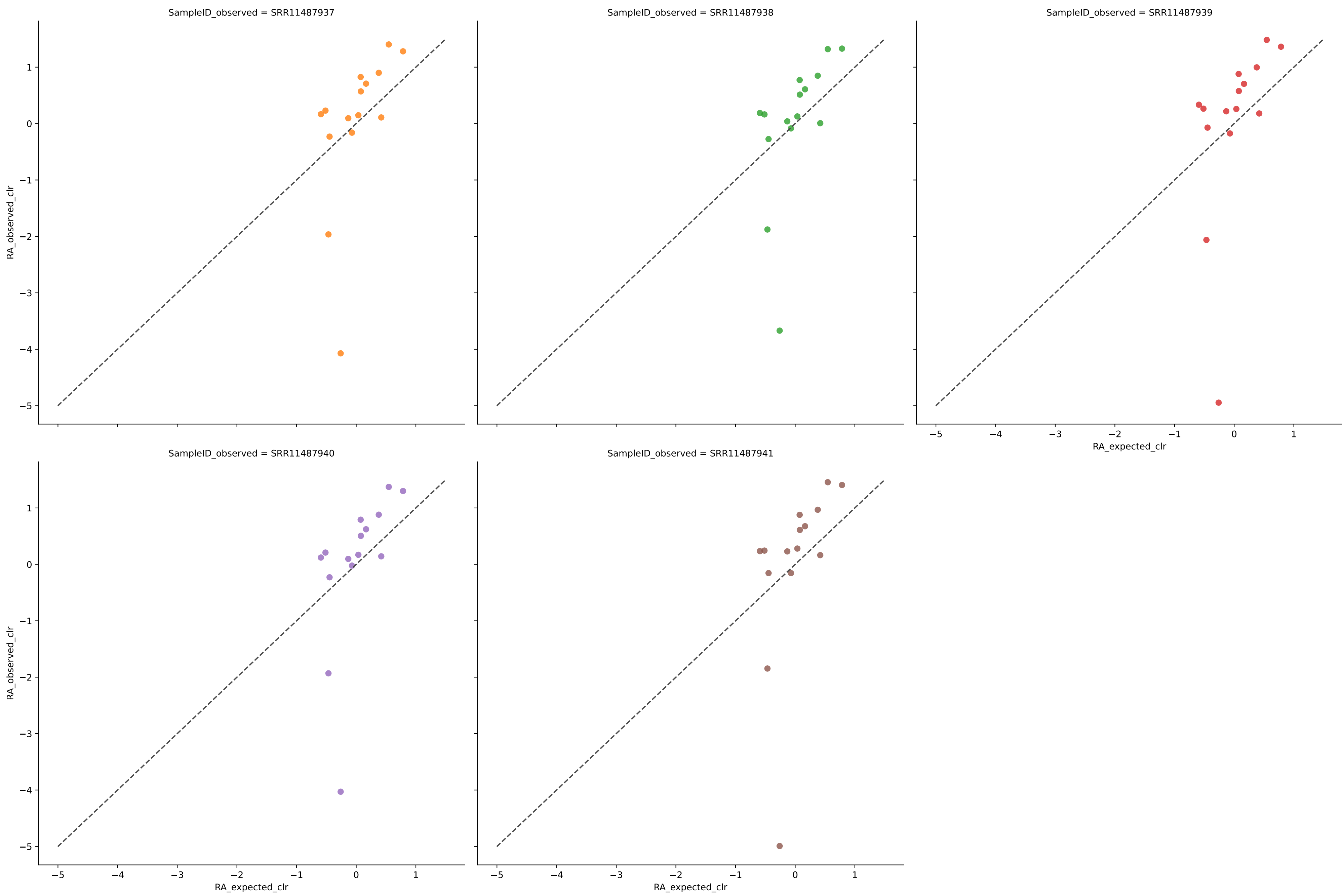
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487931	17	0.9122	0.0044	4.6517	0.8125	0.0084	100.0000	0.0000
SRR11487932	17	0.9108	0.0044	4.7533	0.8110	0.0083	100.0000	0.0000
SRR11487933	17	0.9187	0.0042	4.4721	0.8210	0.0077	100.0000	0.0000
SRR11487934	17	0.9158	0.0043	4.4548	0.8173	0.0079	100.0000	0.0000
SRR11487935	17	0.9223	0.0041	4.4669	0.8255	0.0076	100.0000	0.0000
Average	17	0.9160	0.0043	4.5598	0.8175	0.0080	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using wol in Experiment Amos hilo with filter 1e-05



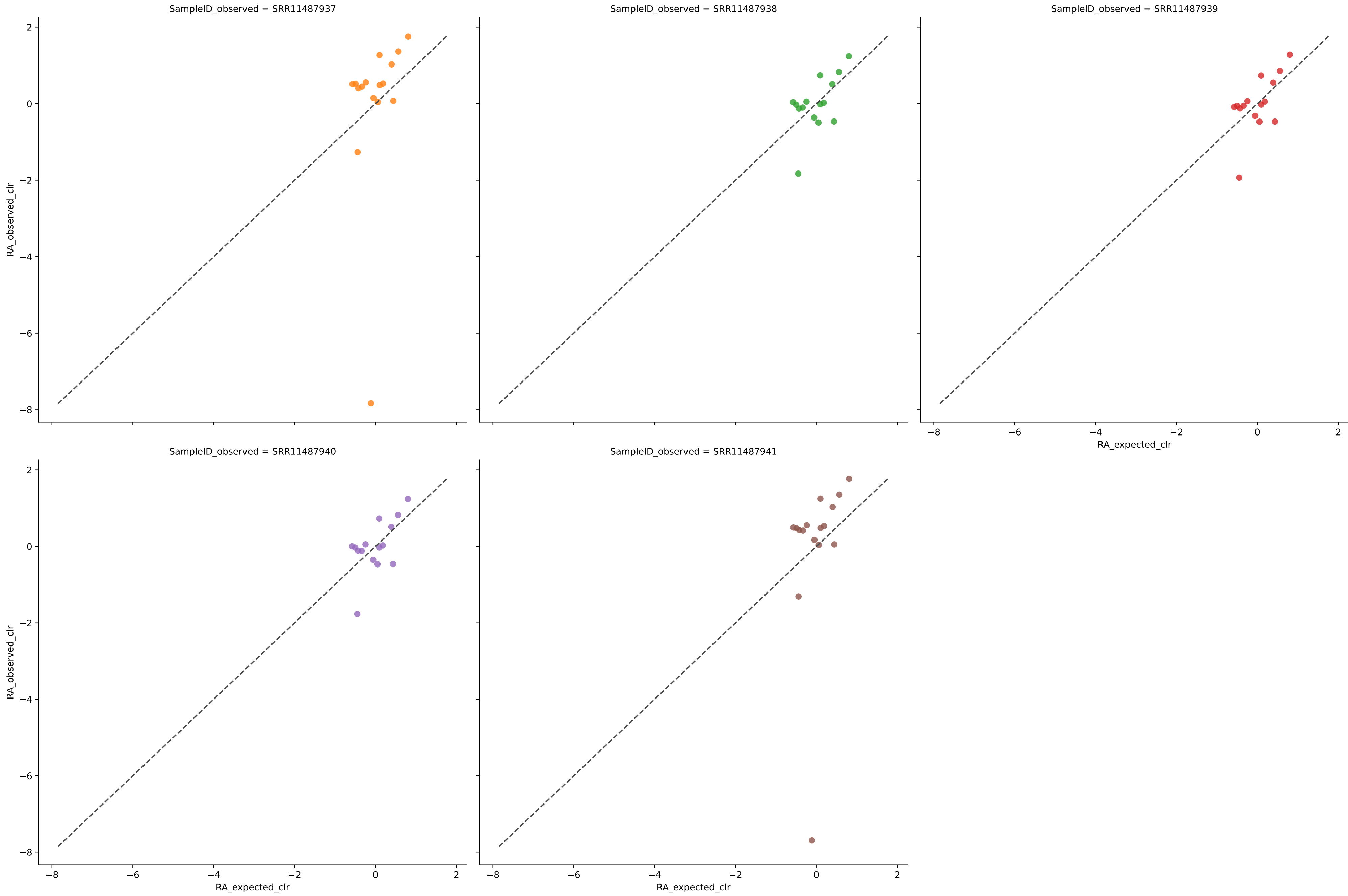
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487931	18	0.2914	0.0083	3.3507	0.6248	0.0177	100.0000	0.0000
SRR11487932	18	0.2756	0.0085	3.4846	0.6160	0.0181	100.0000	0.0000
SRR11487933	18	0.2816	0.0084	3.2745	0.6226	0.0177	100.0000	0.0000
SRR11487934	18	0.2963	0.0083	3.3150	0.6261	0.0175	100.0000	0.0000
SRR11487935	18	0.2883	0.0083	3.2343	0.6285	0.0176	100.0000	0.0000
Average	18	0.2866	0.0084	3.3318	0.6236	0.0177	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using bio4 in Experiment Amos mixed with filter 1e-05



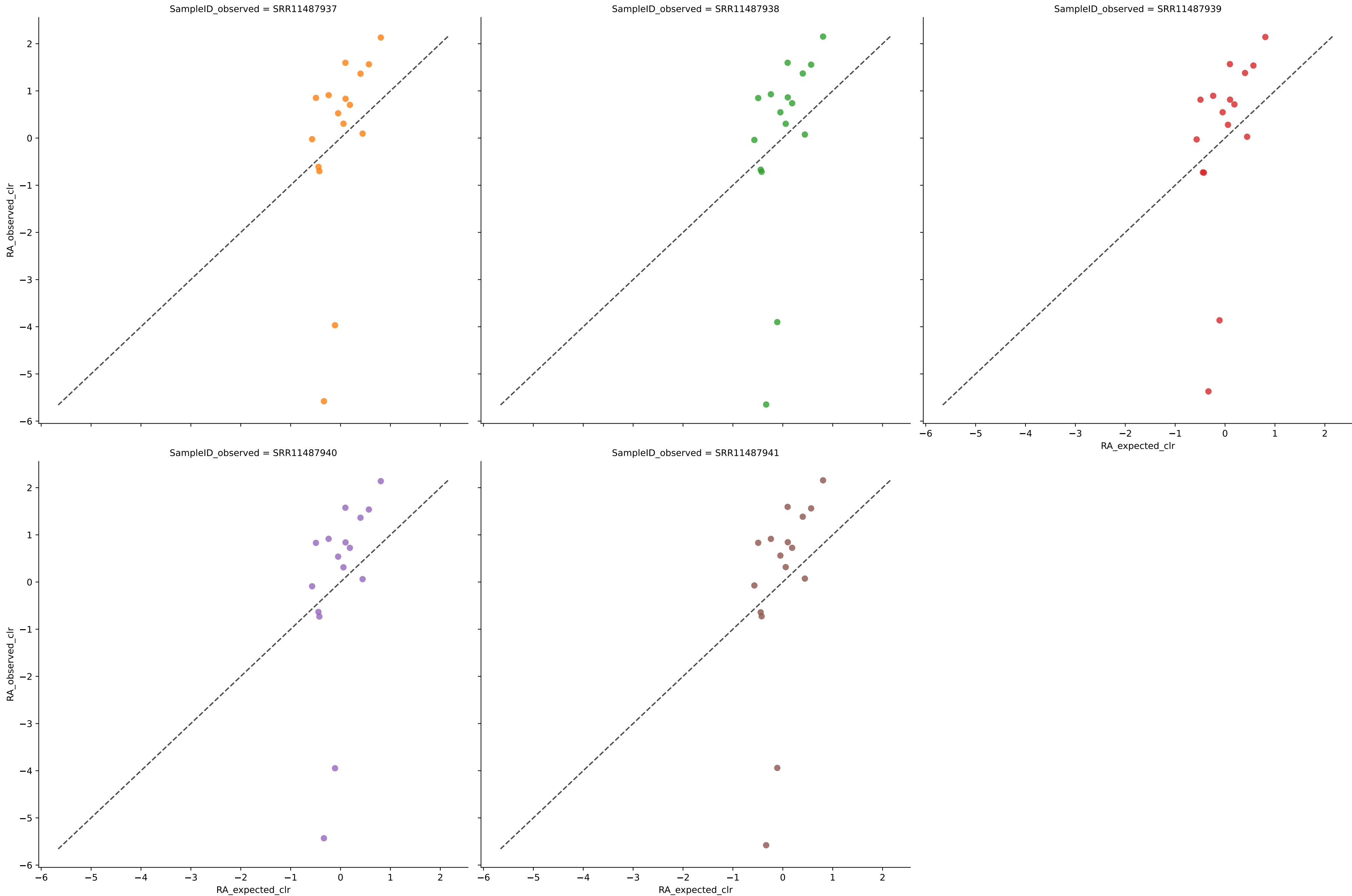
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487937	15	0.6595	0.0047	4.5249	0.8236	0.0058	100.0000	0.0000
SRR11487938	15	0.6856	0.0046	4.1118	0.8260	0.0056	100.0000	0.0000
SRR11487939	15	0.6543	0.0046	5.3975	0.8269	0.0058	100.0000	0.0000
SRR11487940	15	0.6903	0.0044	4.4374	0.8352	0.0055	100.0000	0.0000
SRR11487941	15	0.6856	0.0046	5.3530	0.8283	0.0057	100.0000	0.0000
Average	15	0.6751	0.0046	4.7649	0.8280	0.0057	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using jams in Experiment Amos mixed with filter 1e-05



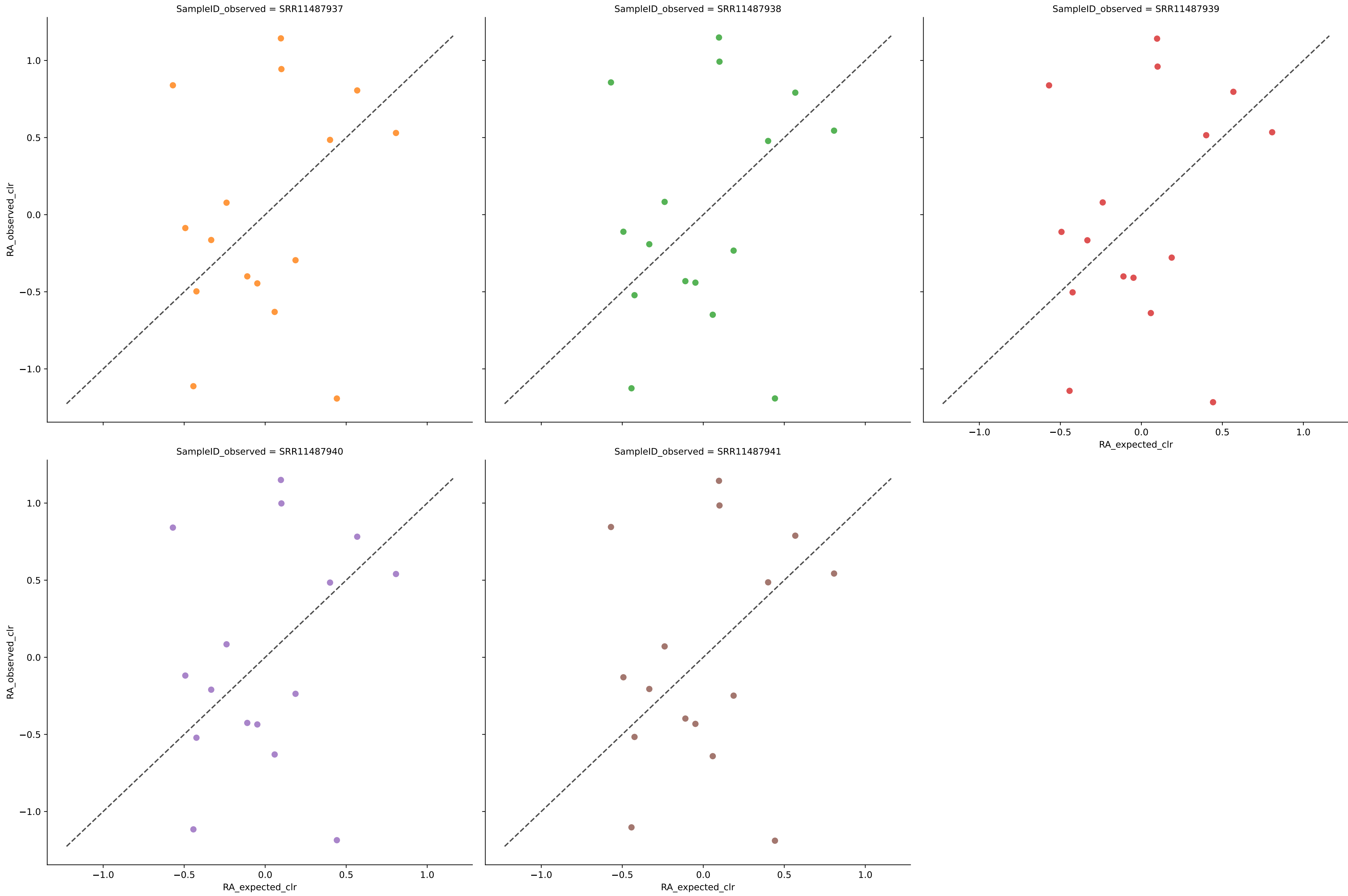
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487937	16	0.5479	0.0051	8.2583	0.7996	0.0062	100.0000	0.0000
SRR11487938	15	0.5697	0.0047	2.1639	0.8210	0.0058	100.0000	0.0000
SRR11487939	15	0.5984	0.0046	2.2036	0.8237	0.0058	100.0000	0.0000
SRR11487940	15	0.5777	0.0047	2.1051	0.8228	0.0058	100.0000	0.0000
SRR11487941	16	0.5581	0.0051	8.1152	0.8006	0.0062	100.0000	0.0000
Average	15	0.5704	0.0048	4.5692	0.8135	0.0059	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using wgsa in Experiment Amos mixed with filter 1e-05



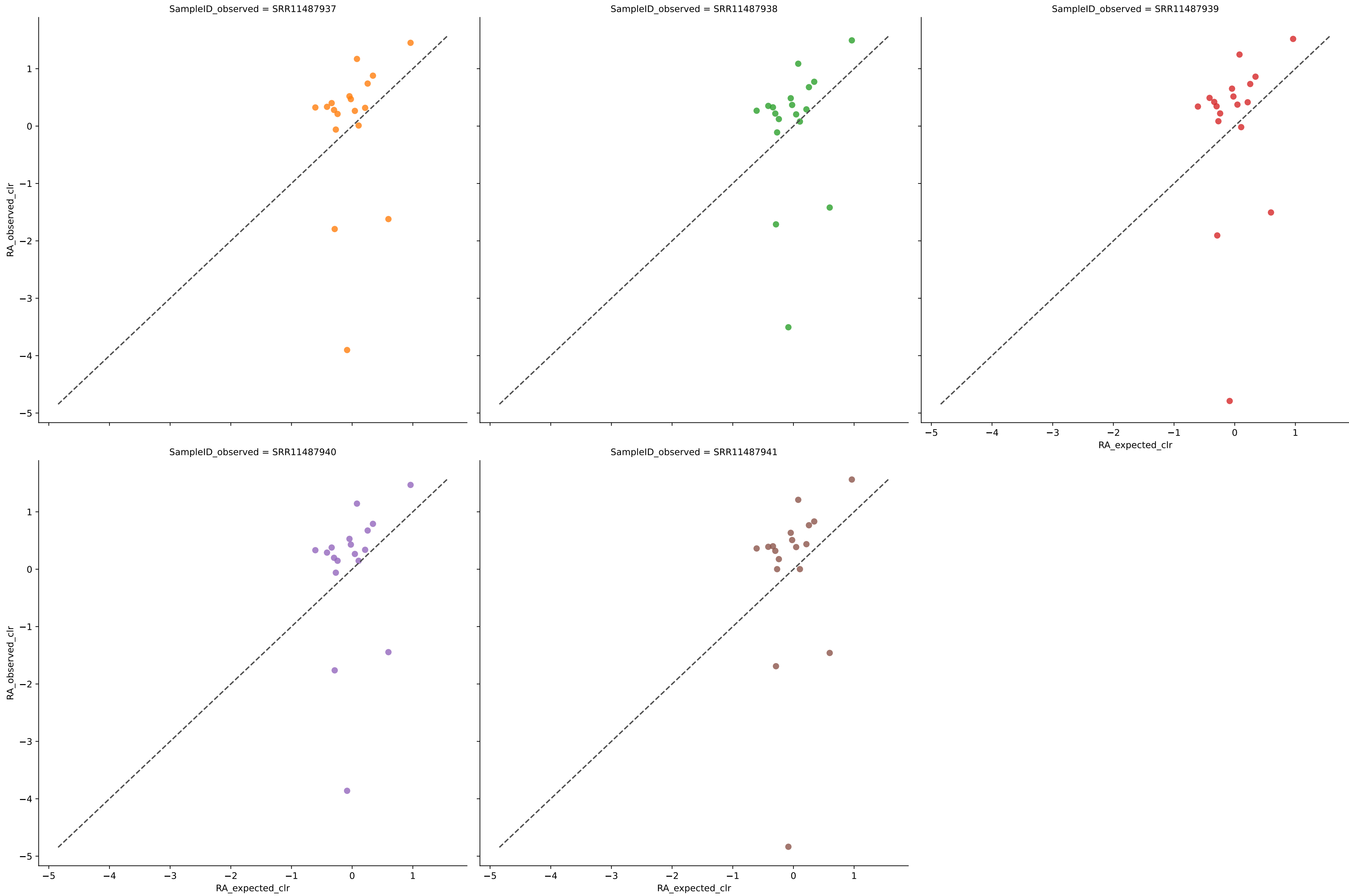
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487937	16	0.6162	0.0064	7.2906	0.7422	0.0081	100.0000	0.0000
SRR11487938	16	0.6172	0.0064	7.3210	0.7424	0.0082	100.0000	0.0000
SRR11487939	16	0.6217	0.0065	7.0792	0.7405	0.0082	100.0000	0.0000
SRR11487940	16	0.6200	0.0064	7.1672	0.7428	0.0081	100.0000	0.0000
SRR11487941	16	0.6231	0.0065	7.2852	0.7413	0.0082	100.0000	0.0000
Average	16	0.6196	0.0065	7.2286	0.7418	0.0082	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using wol in Experiment Amos mixed with filter 1e-05



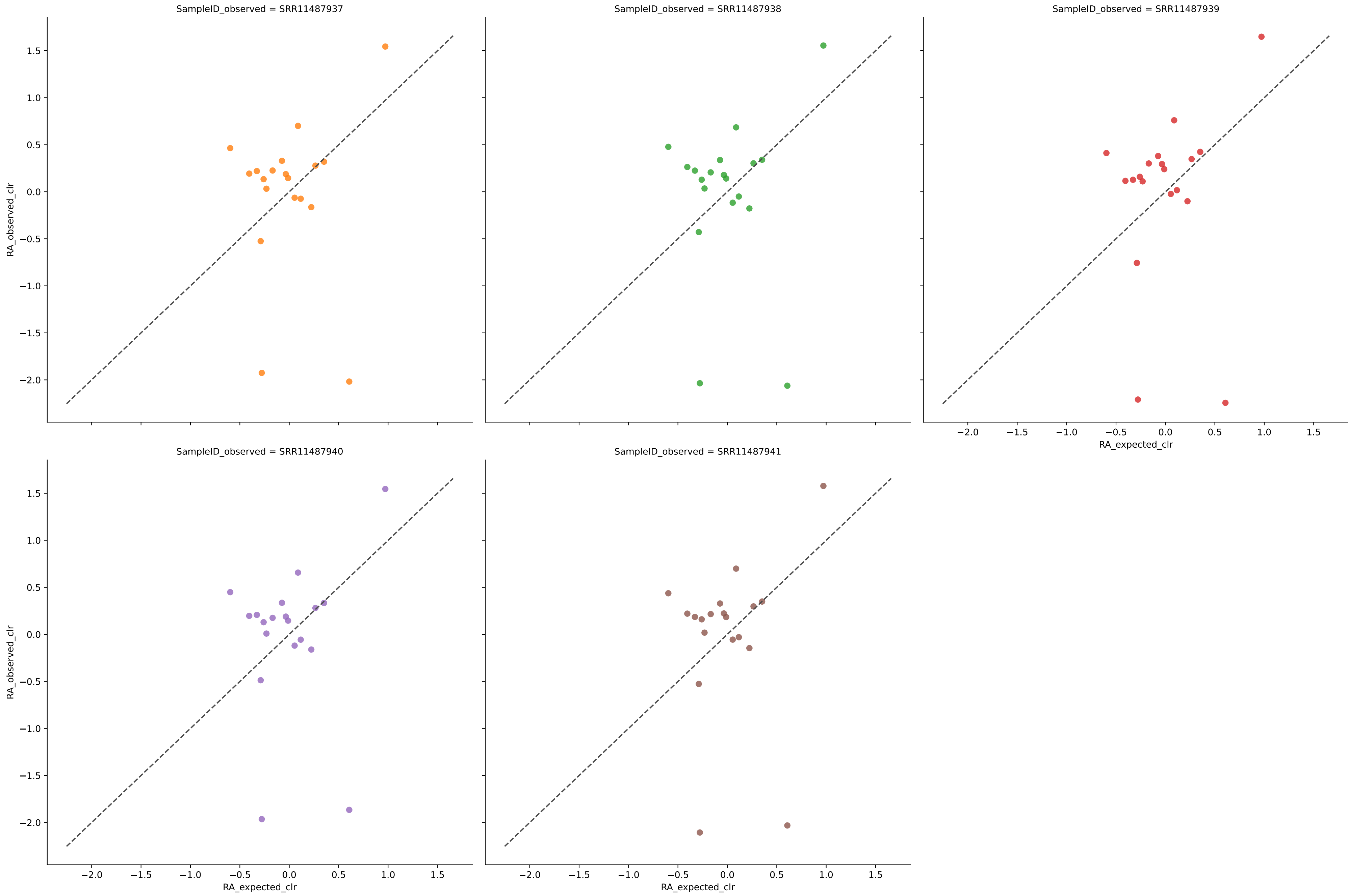
	Diversity	R ²	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487937	16	0.0645	0.0066	2.8809	0.7364	0.0087	100.0000	0.0000
SRR11487938	16	0.0633	0.0066	2.9006	0.7340	0.0089	100.0000	0.0000
SRR11487939	16	0.0662	0.0065	2.8965	0.7387	0.0087	100.0000	0.0000
SRR11487940	16	0.0643	0.0066	2.8819	0.7351	0.0088	100.0000	0.0000
SRR11487941	16	0.0658	0.0066	2.8748	0.7369	0.0088	100.0000	0.0000
Average	16	0.0648	0.0066	2.8869	0.7362	0.0088	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using bio4 in Experiment Amos mixed with filter 1e-05



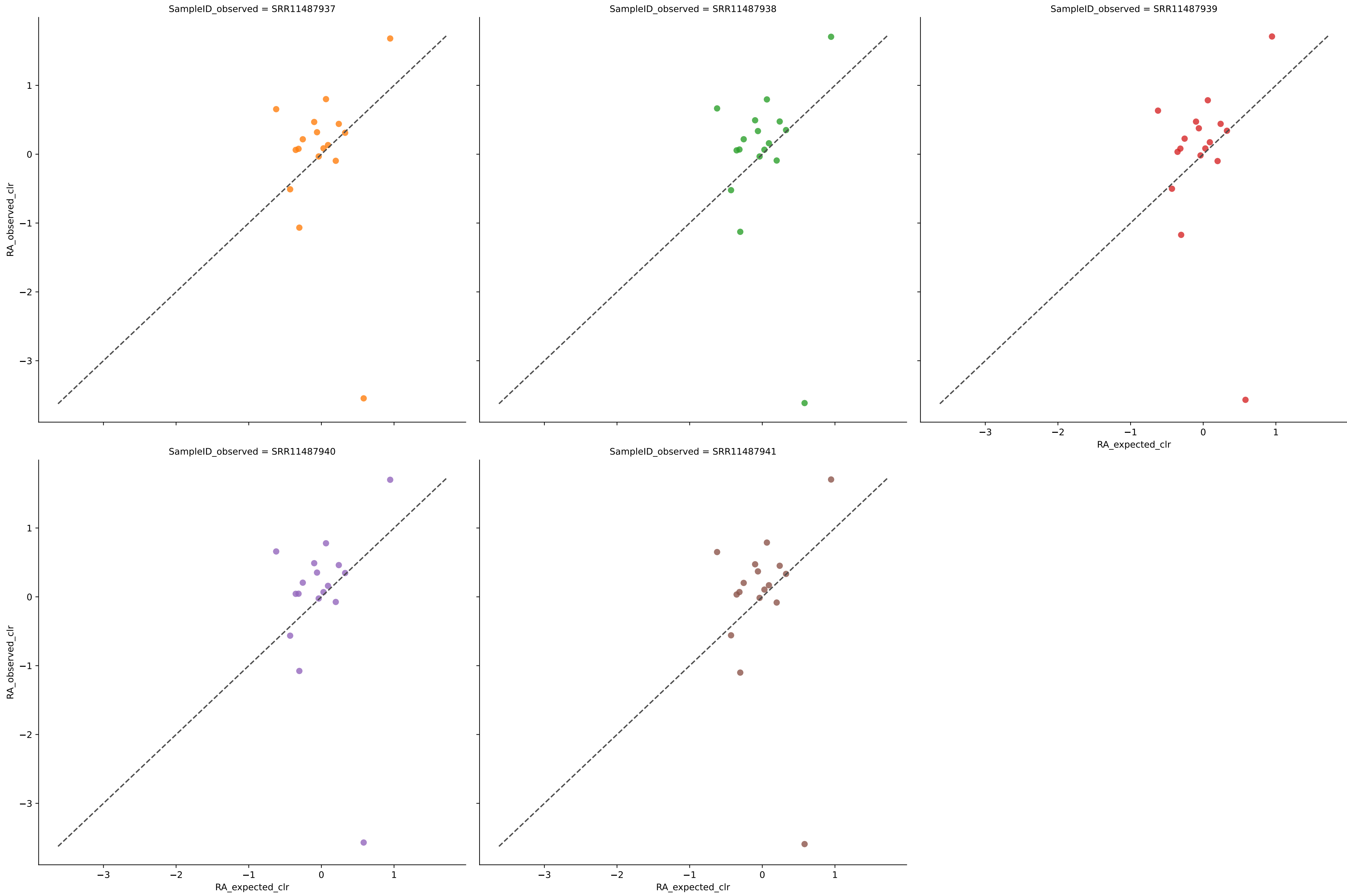
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487937	18	0.2996	0.0046	5.1875	0.7914	0.0063	100.0000	0.0000
SRR11487938	18	0.3547	0.0045	4.7066	0.7969	0.0062	100.0000	0.0000
SRR11487939	18	0.2928	0.0046	5.9450	0.7951	0.0063	100.0000	0.0000
SRR11487940	18	0.3277	0.0044	5.0307	0.8030	0.0061	100.0000	0.0000
SRR11487941	18	0.3332	0.0045	5.8823	0.7971	0.0063	100.0000	0.0000
Average	18	0.3216	0.0045	5.3504	0.7967	0.0062	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using jams in Experiment Amos mixed with filter 1e-05



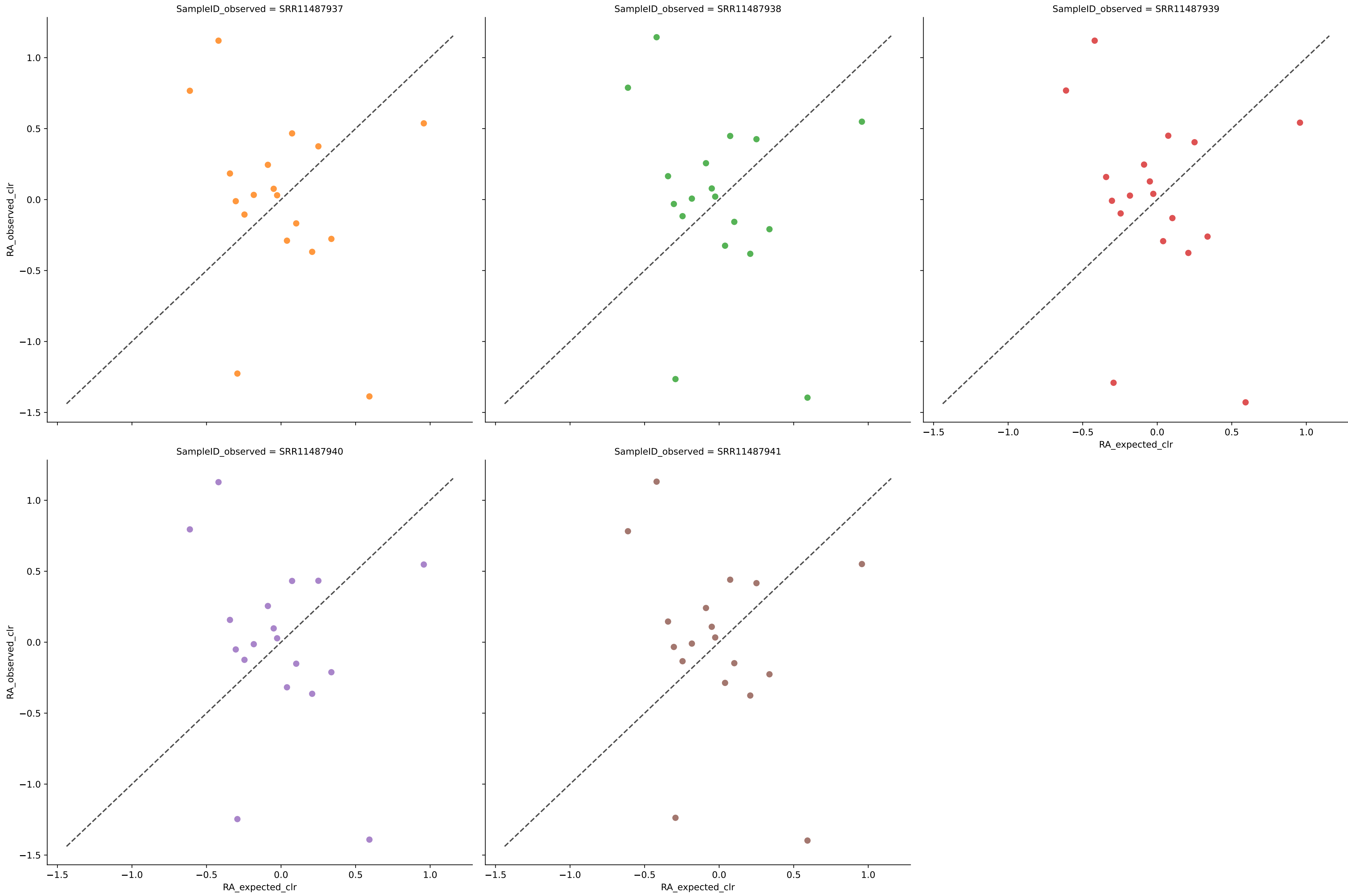
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487937	19	0.3708	0.0043	3.6003	0.7947	0.0060	100.0000	0.0000
SRR11487938	19	0.3682	0.0044	3.6950	0.7941	0.0061	100.0000	0.0000
SRR11487939	19	0.4162	0.0042	3.9369	0.7974	0.0060	100.0000	0.0000
SRR11487940	19	0.3834	0.0043	3.4885	0.7968	0.0060	100.0000	0.0000
SRR11487941	19	0.3887	0.0043	3.6944	0.7964	0.0060	100.0000	0.0000
Average	19	0.3854	0.0043	3.6830	0.7959	0.0060	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using wgsa in Experiment Amos mixed with filter 1e-05



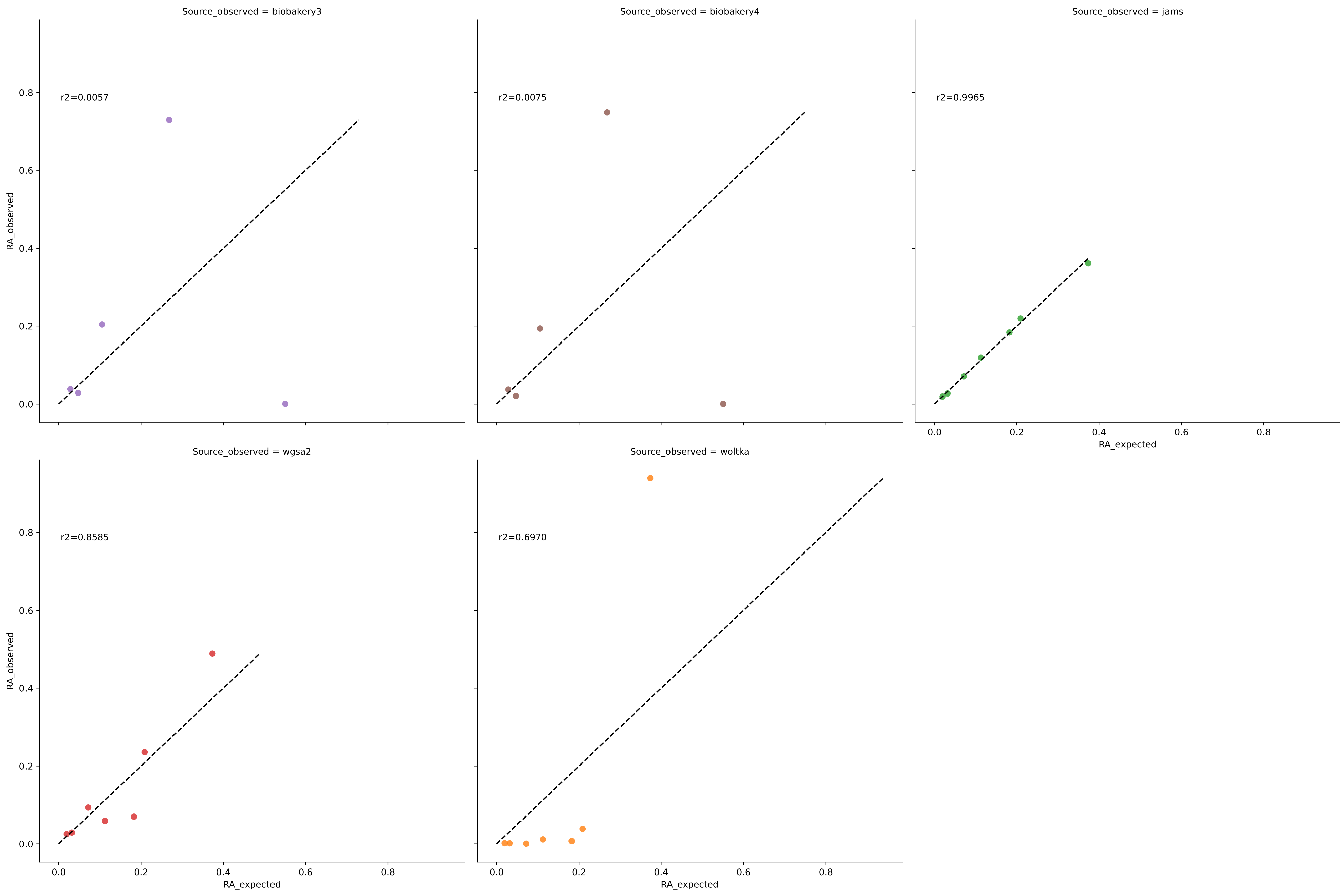
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487937	17	0.3833	0.0051	4.6362	0.7831	0.0074	100.0000	0.0000
SRR11487938	17	0.3924	0.0051	4.7186	0.7829	0.0074	100.0000	0.0000
SRR11487939	17	0.3983	0.0051	4.6751	0.7827	0.0075	100.0000	0.0000
SRR11487940	17	0.3953	0.0051	4.6599	0.7843	0.0074	100.0000	0.0000
SRR11487941	17	0.3961	0.0051	4.6848	0.7841	0.0074	100.0000	0.0000
Average	17	0.3931	0.0051	4.6749	0.7834	0.0074	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using wol in Experiment Amos mixed with filter 1e-05

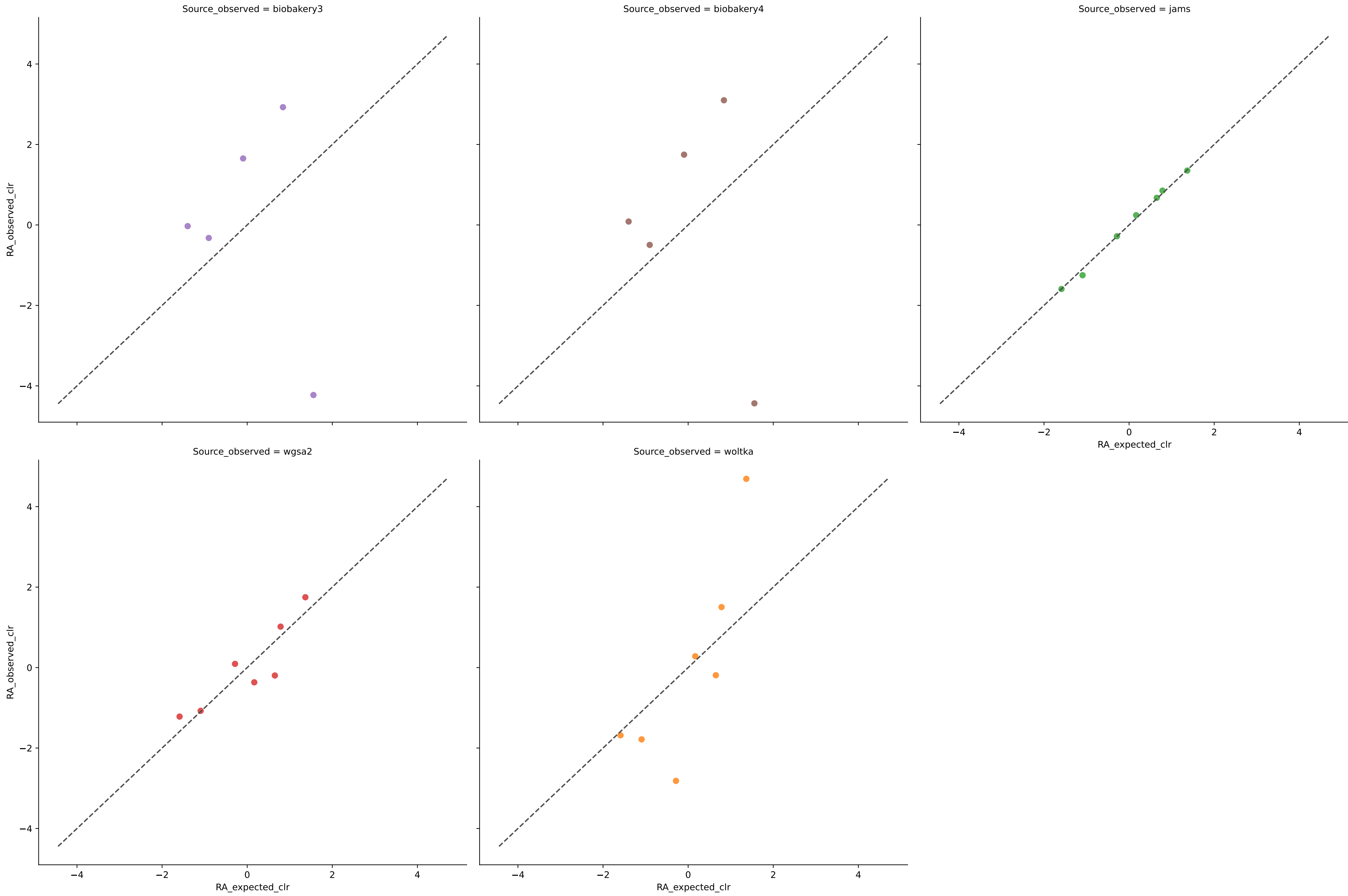


	Diversity	R ²	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487937	18	0.0234	0.0058	3.2948	0.7376	0.0085	100.0000	0.0000
SRR11487938	18	0.0204	0.0058	3.3182	0.7373	0.0085	100.0000	0.0000
SRR11487939	18	0.0221	0.0058	3.3334	0.7377	0.0084	100.0000	0.0000
SRR11487940	18	0.0196	0.0058	3.2964	0.7397	0.0084	100.0000	0.0000
SRR11487941	18	0.0193	0.0058	3.2935	0.7407	0.0084	100.0000	0.0000
Average	18	0.0210	0.0058	3.3073	0.7386	0.0085	100.0000	0.0000

Bivariate Linear Regression for Sample S1 in Experiment bmock12 (Genus at filter threshold 0.0001)

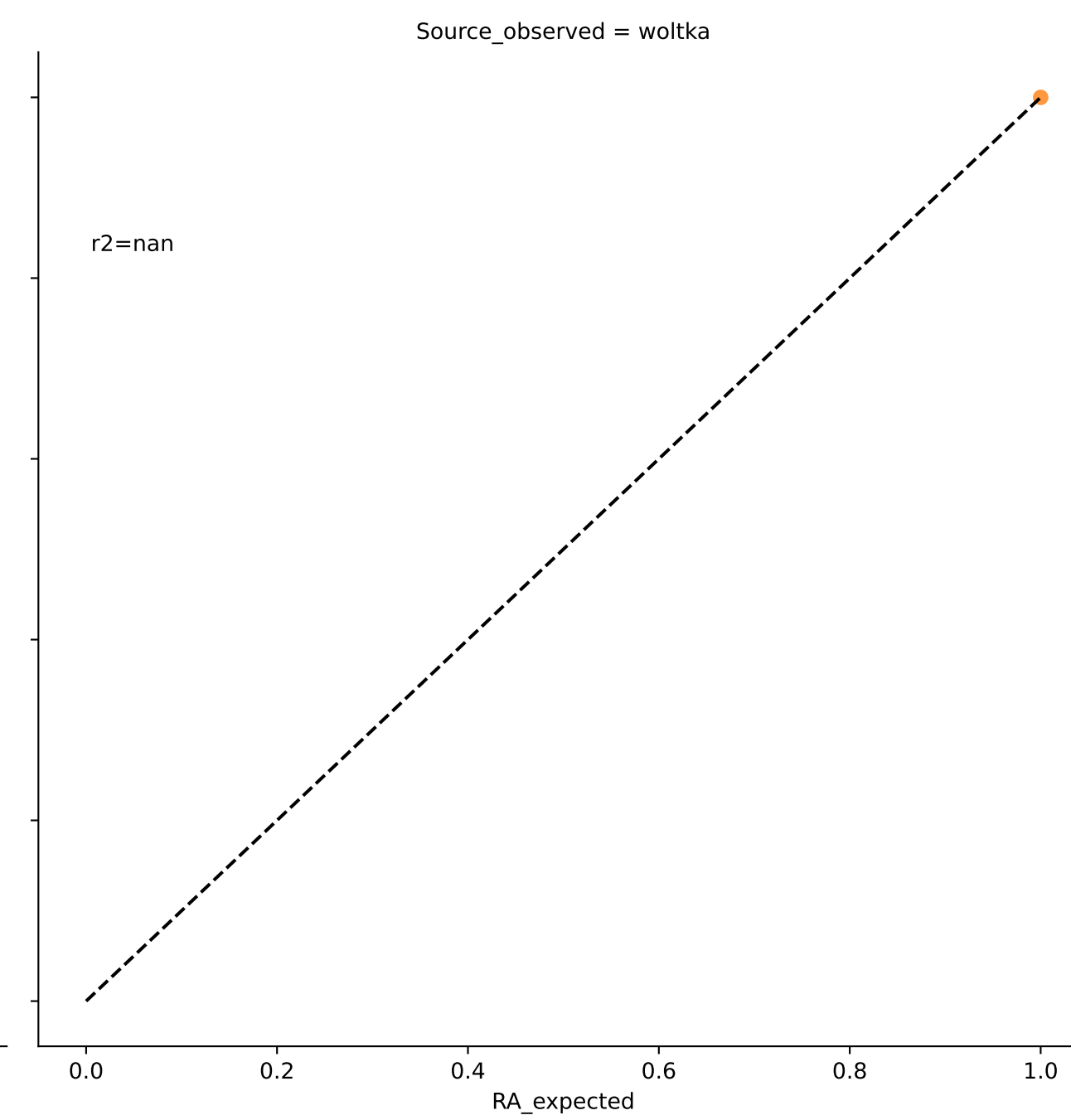
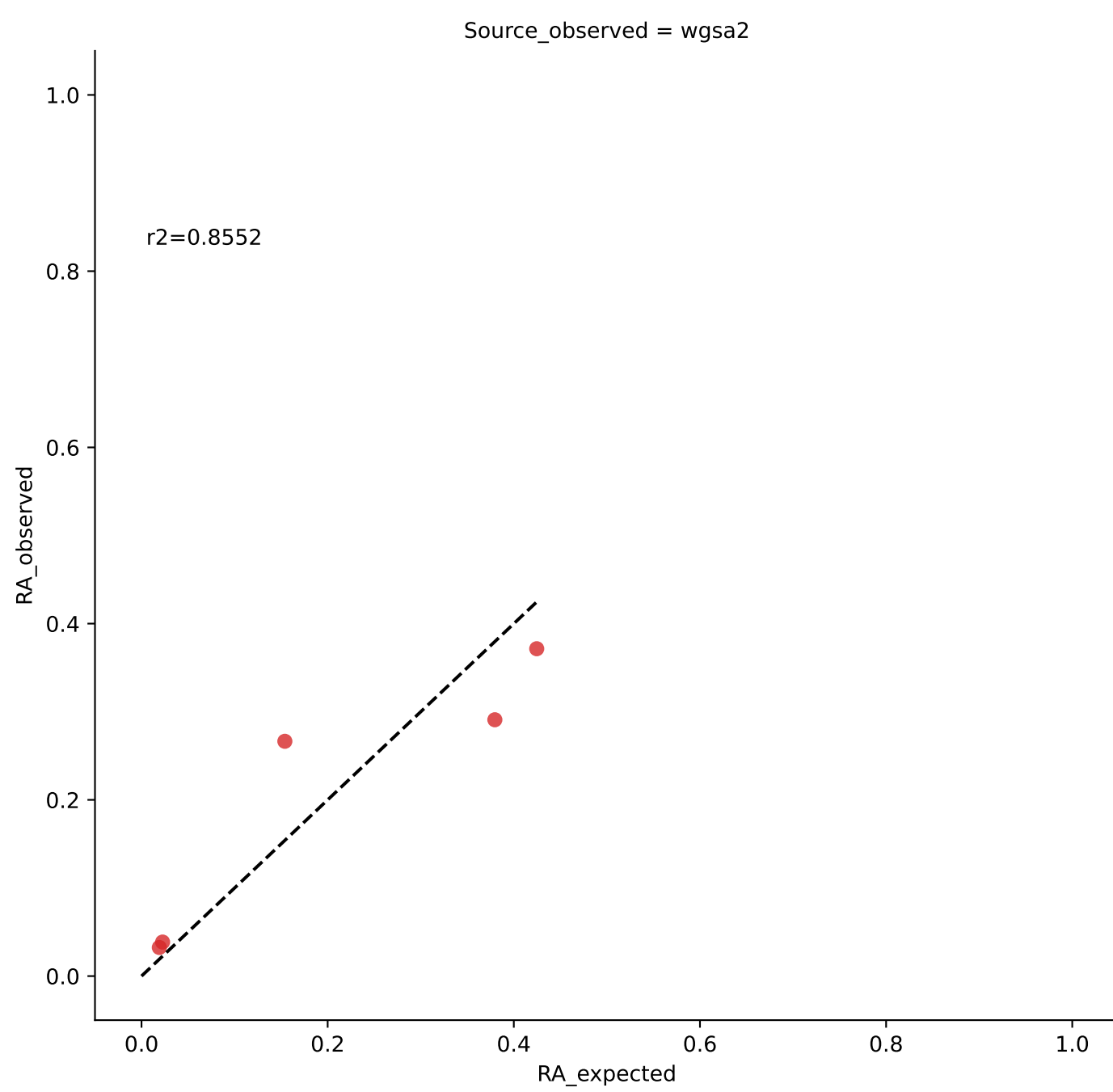
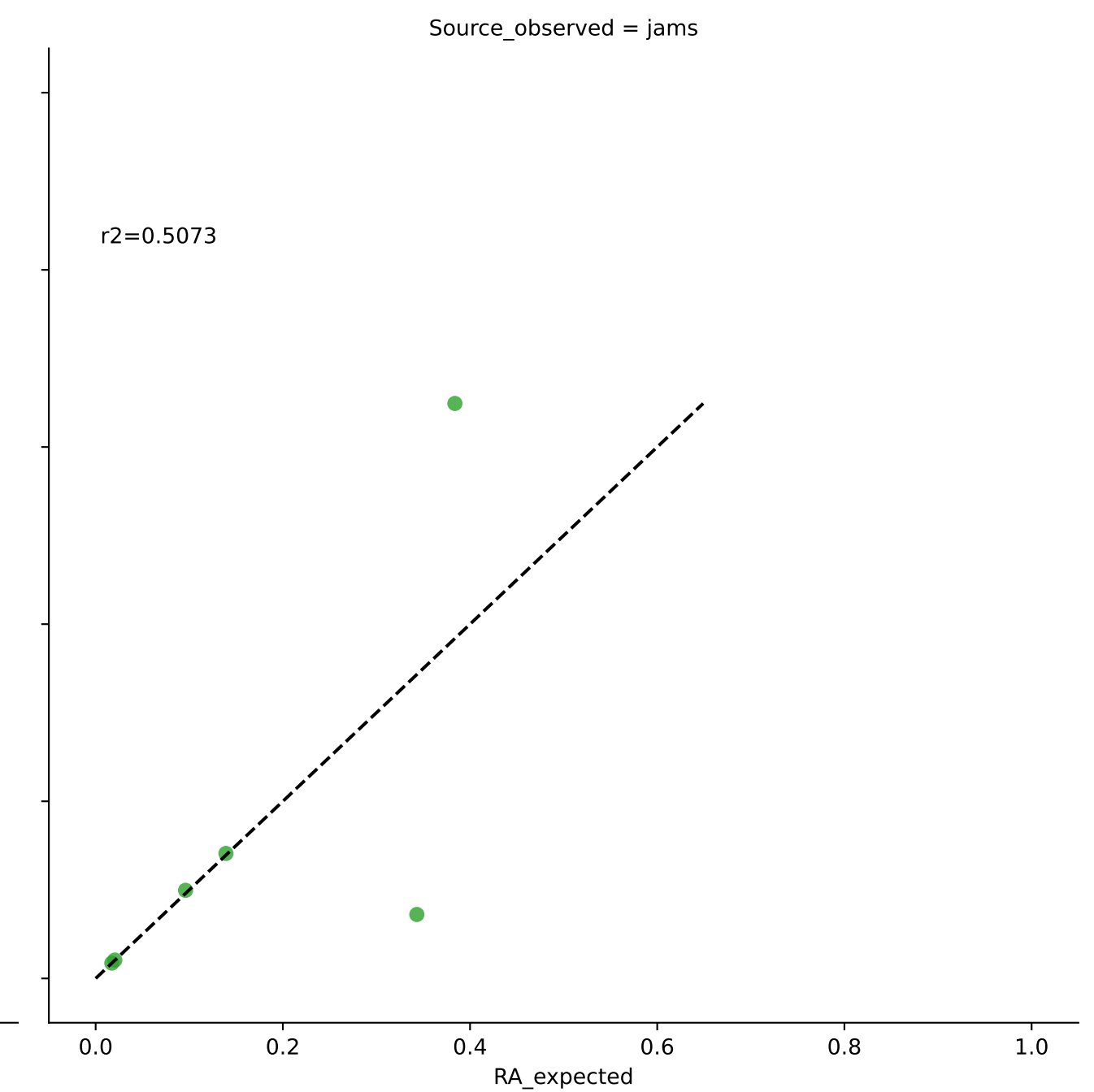
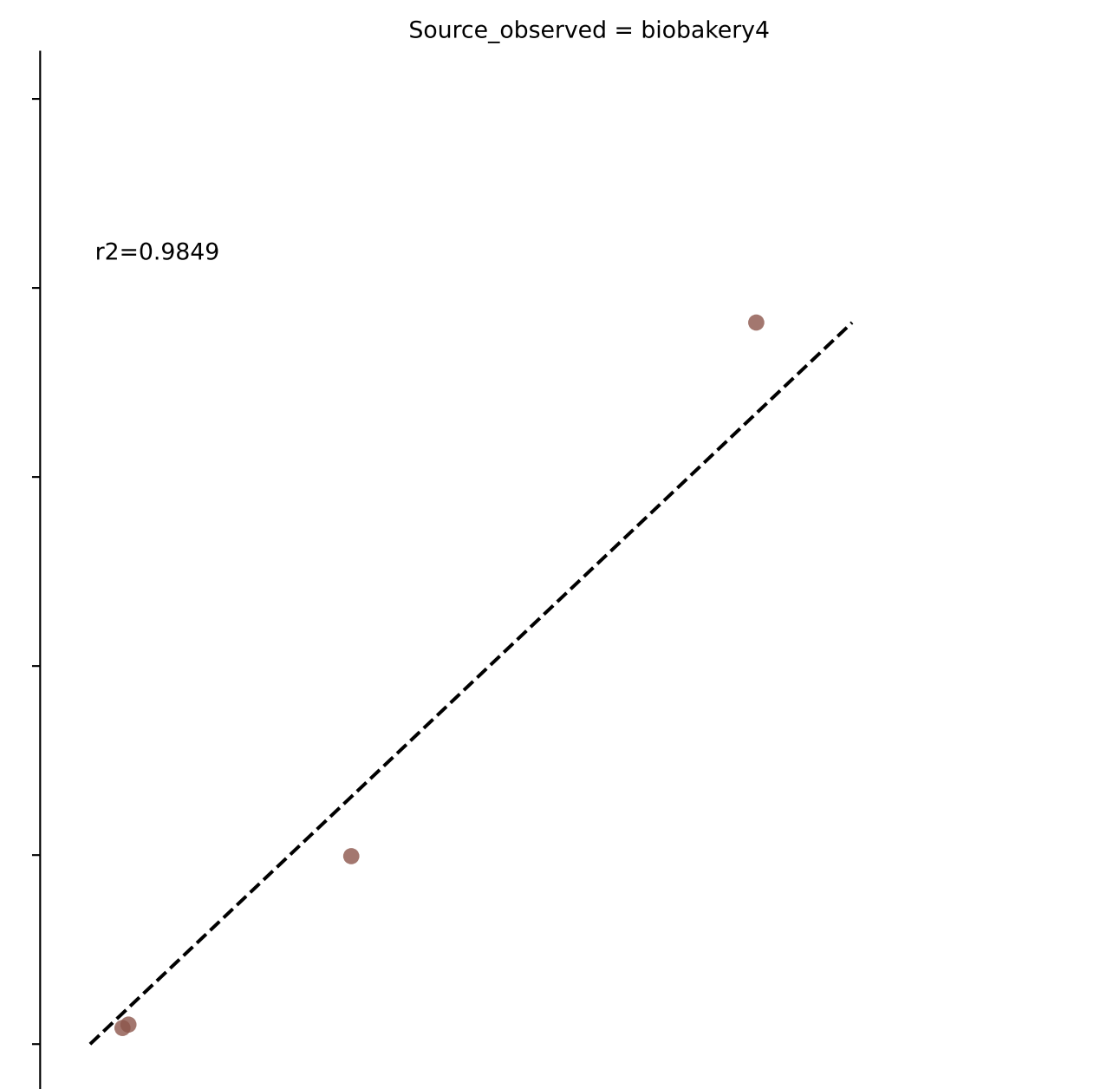
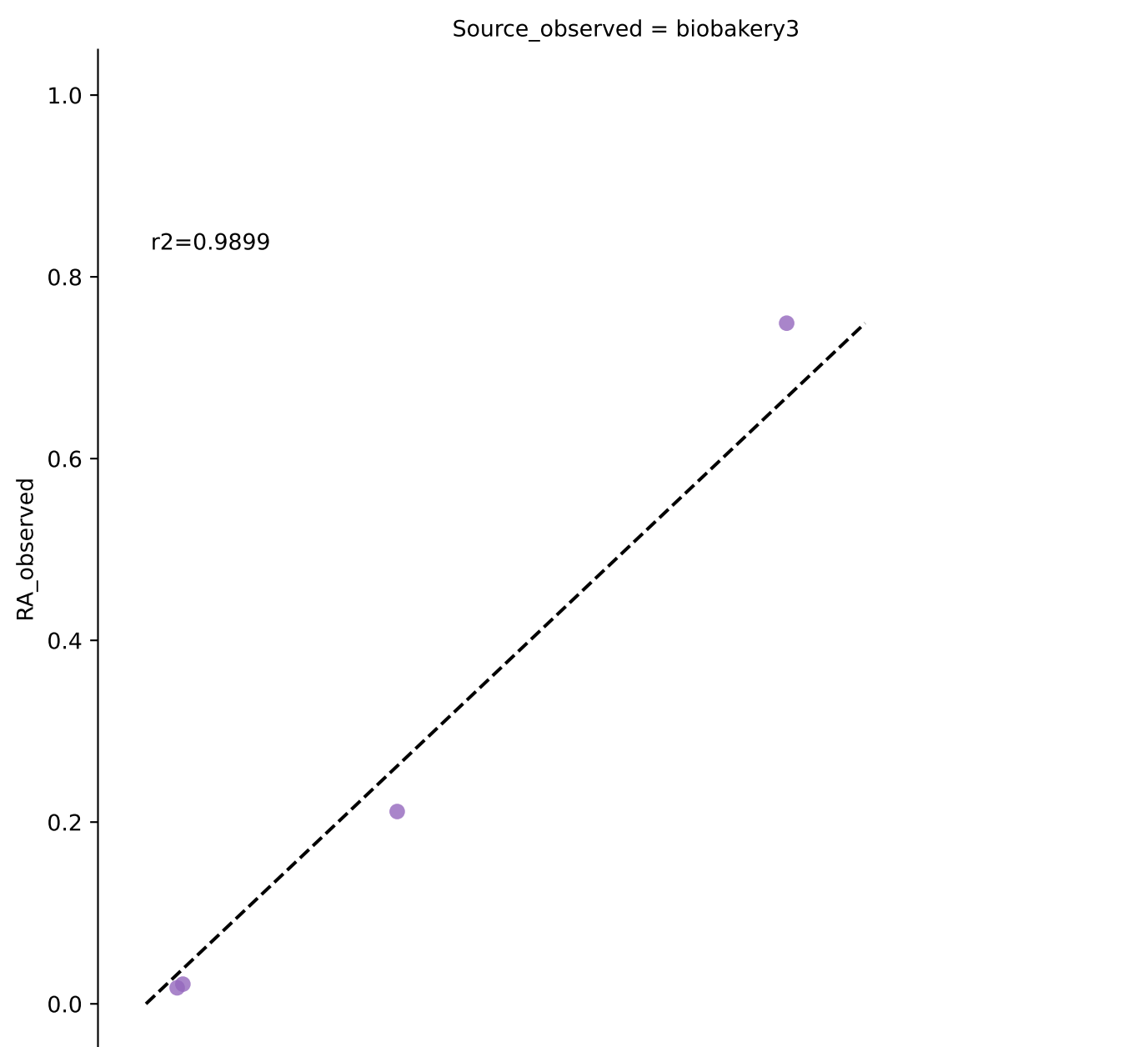


Bivariate Linear Regression for Sample S1 in Experiment bmock12 (Genus at filter threshold 0.0001)

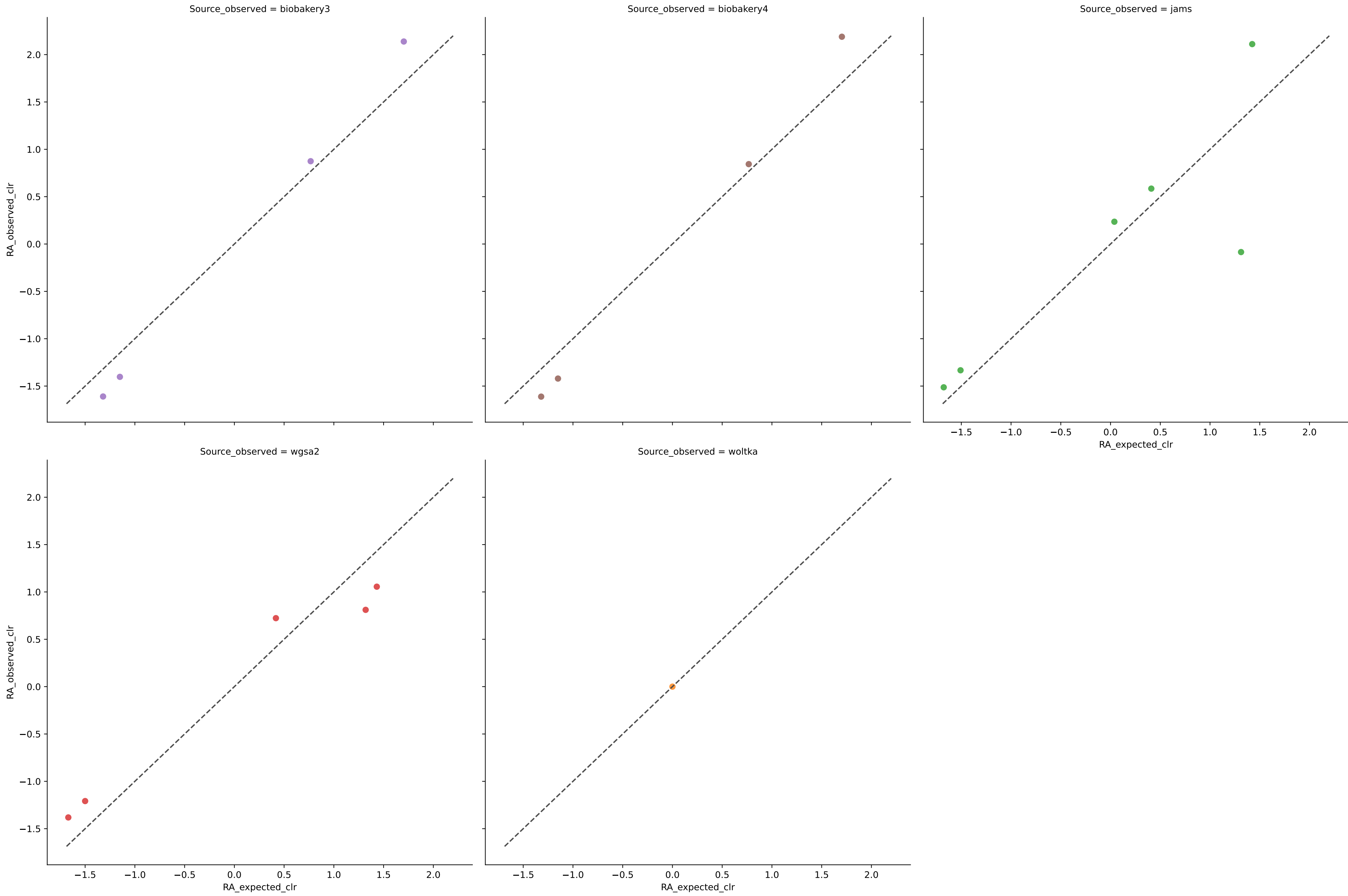


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
biobakery3	5	0.0057	0.2274	6.5633	0.4316	0.3238	100.0000	0.0000
biobakery4	5	0.0075	0.2305	6.8378	0.4237	0.3290	100.0000	0.0000
jams	7	0.9965	0.0053	0.1890	0.9813	0.0070	100.0000	0.0000
wgsa2	7	0.8585	0.0483	1.2171	0.8311	0.0654	100.0000	0.0000
woltka	7	0.6970	0.1616	4.3790	0.4344	0.2379	100.0000	0.0000

Bivariate Linear Regression for Sample S1 in Experiment bmock12 (Species at filter threshold 0.0001)

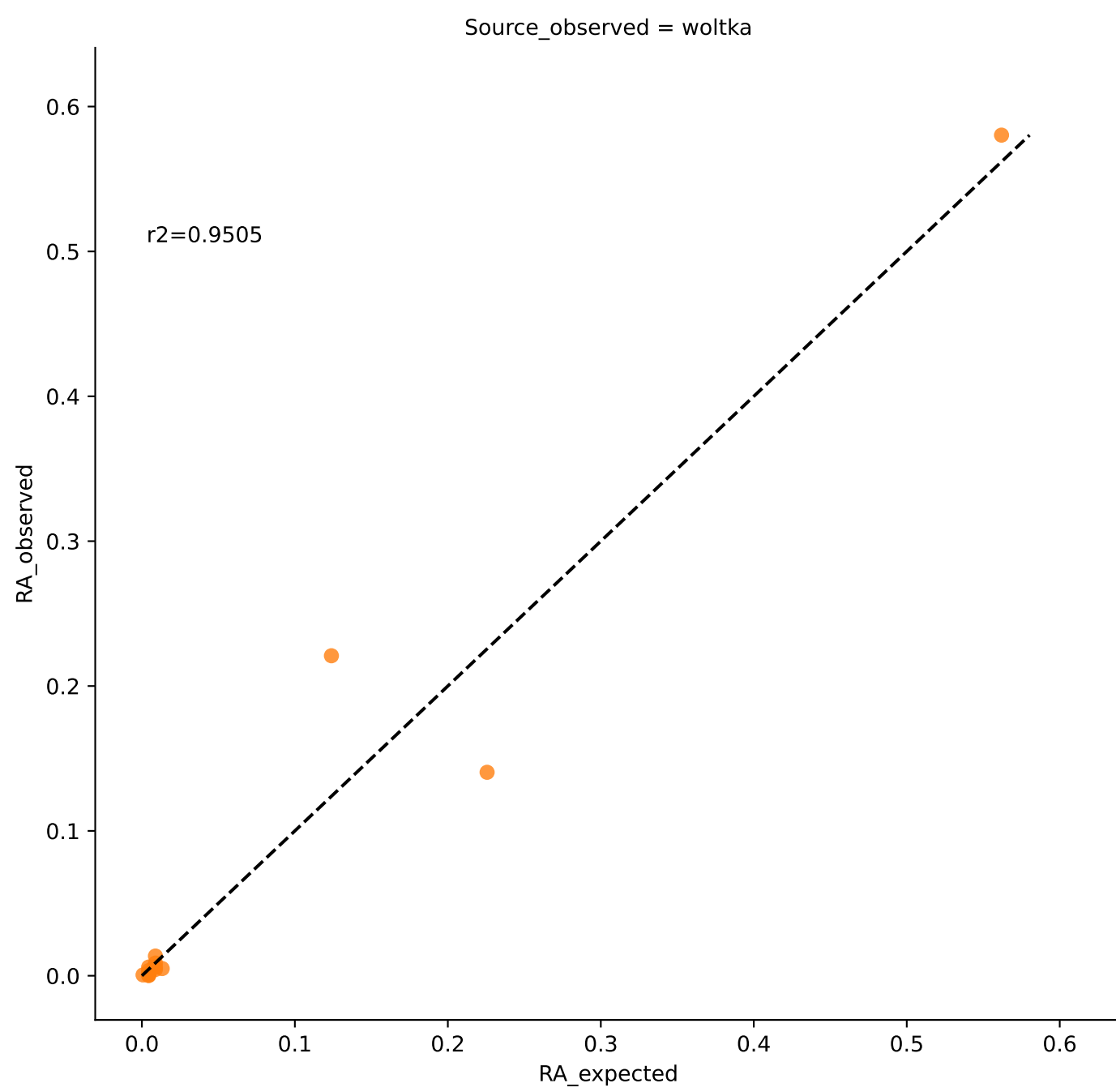
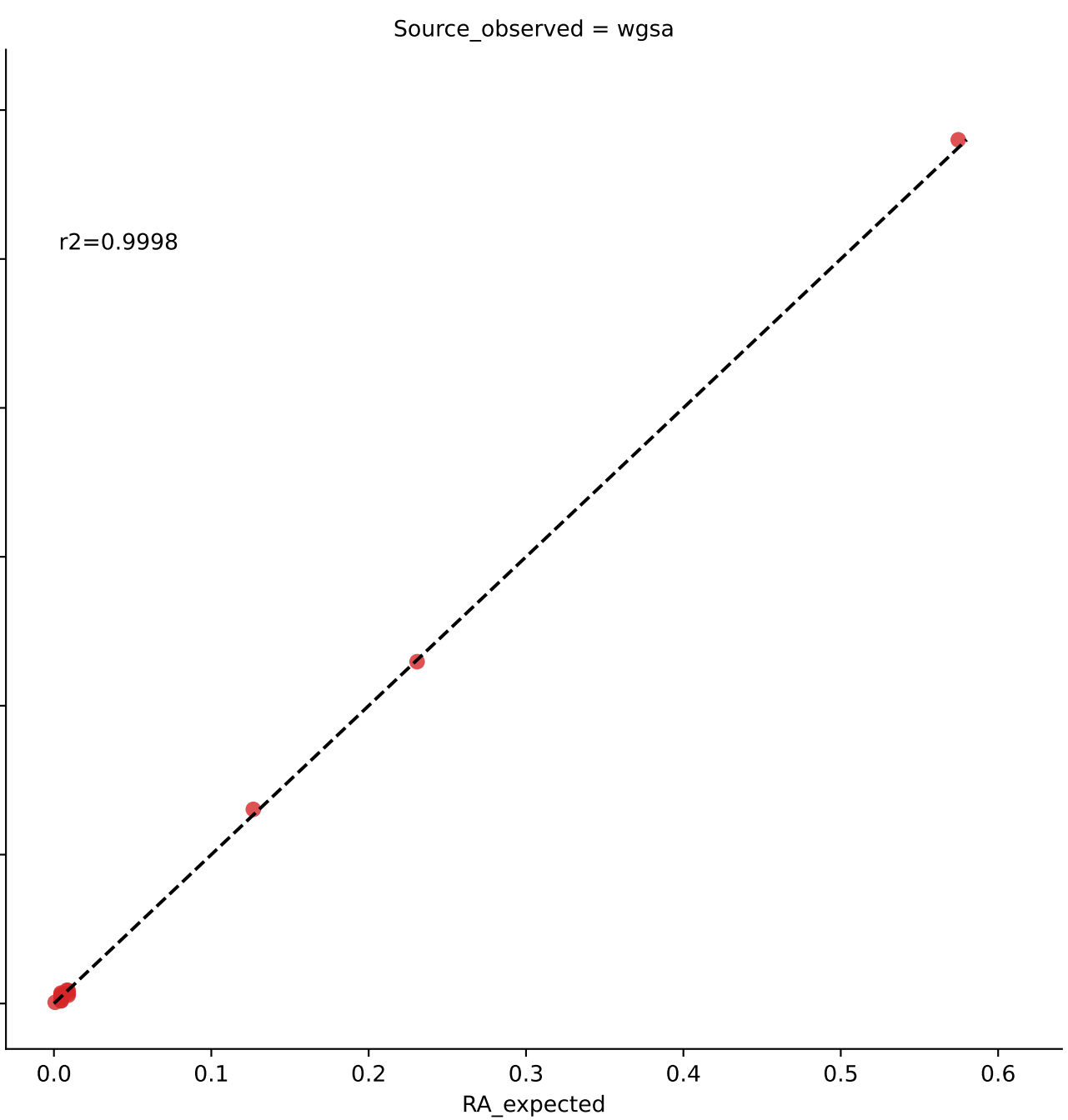
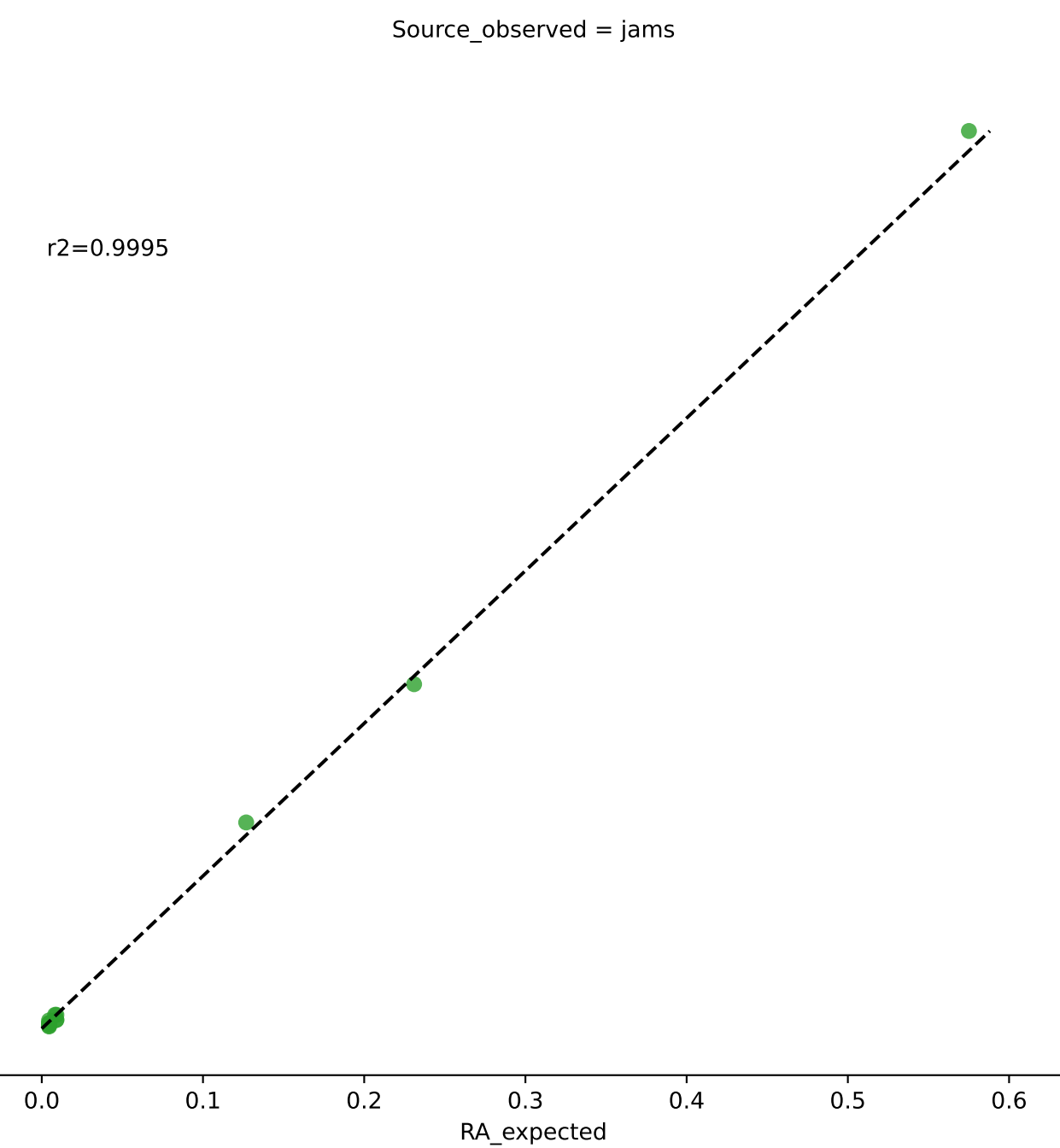
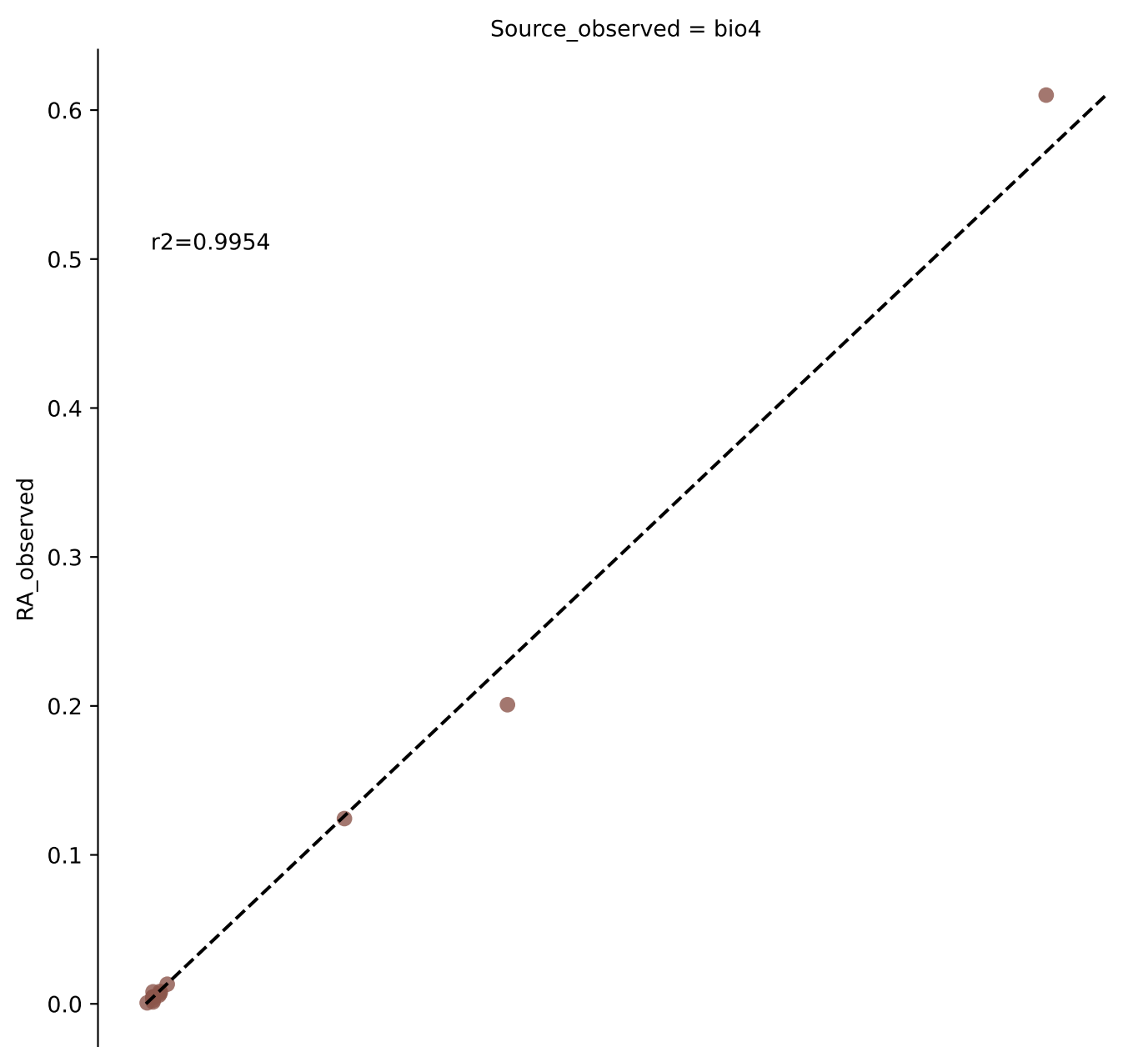


Bivariate Linear Regression for Sample S1 in Experiment bmock12 (Species at filter threshold 0.0001)

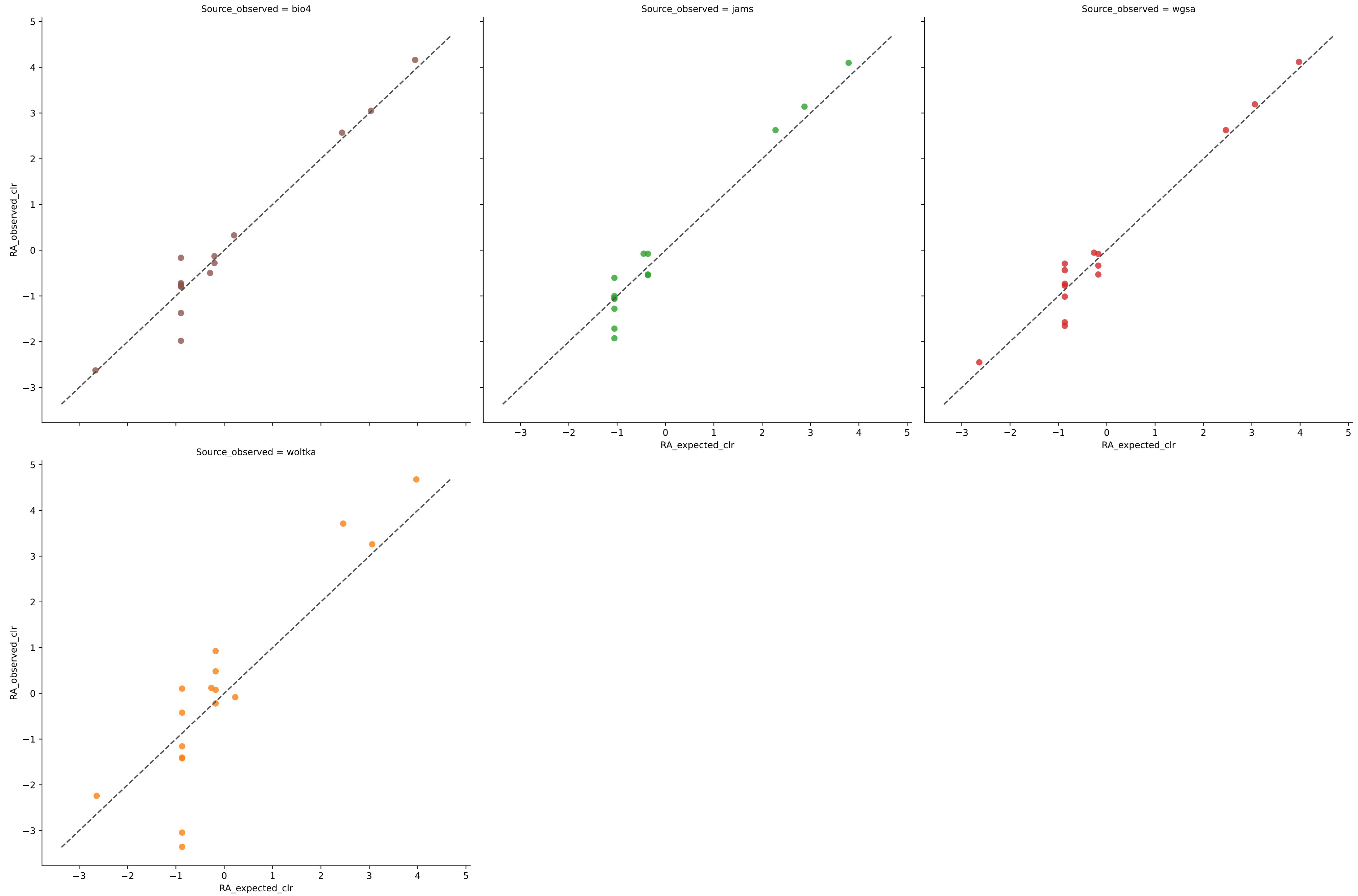


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
biobakery3	4	0.9899	0.0408	0.5907	0.9184	0.0492	100.0000	0.0000
biobakery4	4	0.9849	0.0480	0.6324	0.9040	0.0586	100.0000	0.0000
jams	6	0.5073	0.0903	1.5970	0.7290	0.1548	100.0000	0.0000
wgsa2	5	0.8552	0.0567	0.8135	0.8582	0.0690	100.0000	0.0000
woltka	1	nan	0.0000	0.0000	1.0000	0.0000	100.0000	0.0000

Bivariate Linear Regression for Sample S1 in Experiment camisimGI (Genus at filter threshold 0.0001)

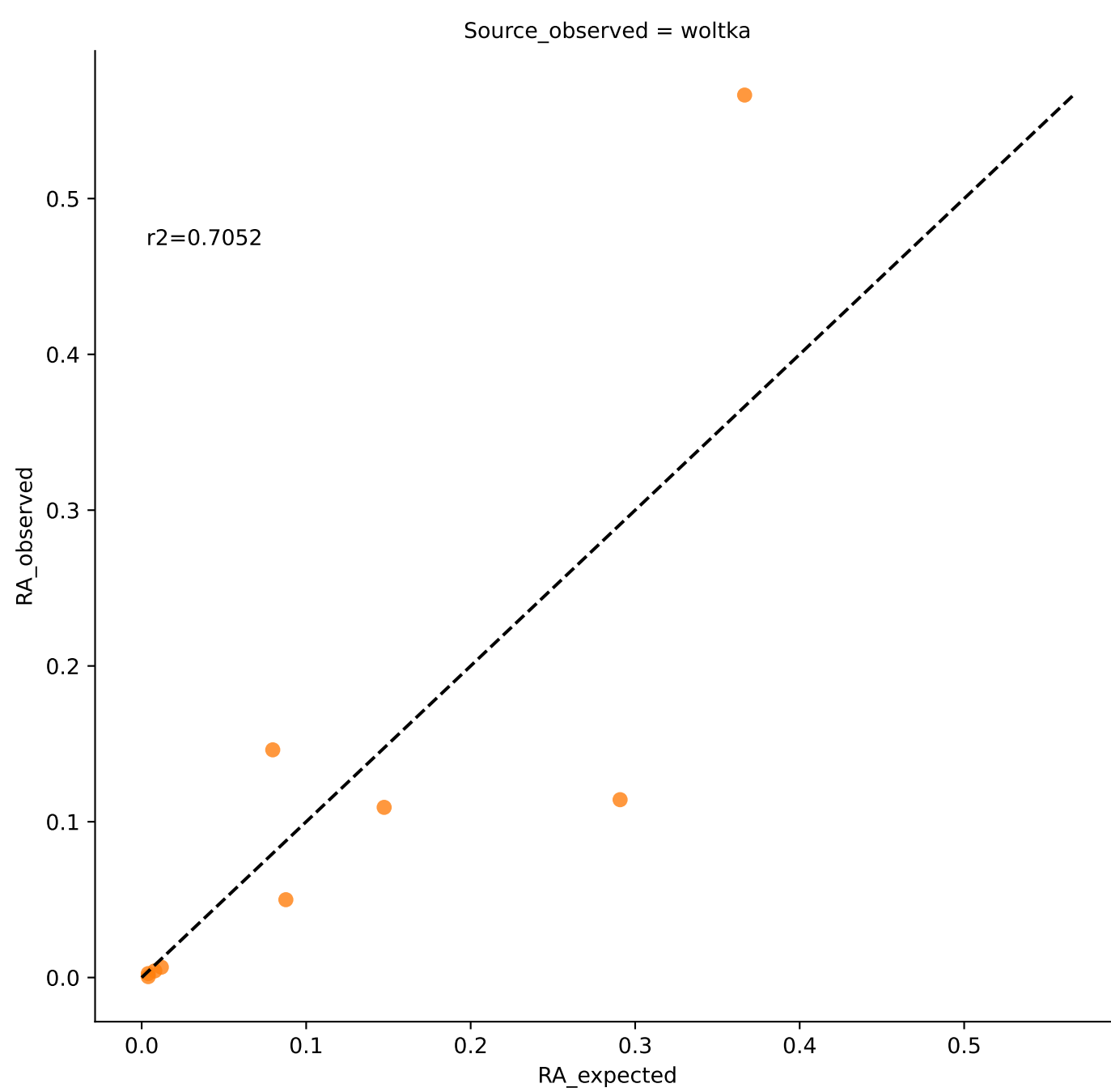
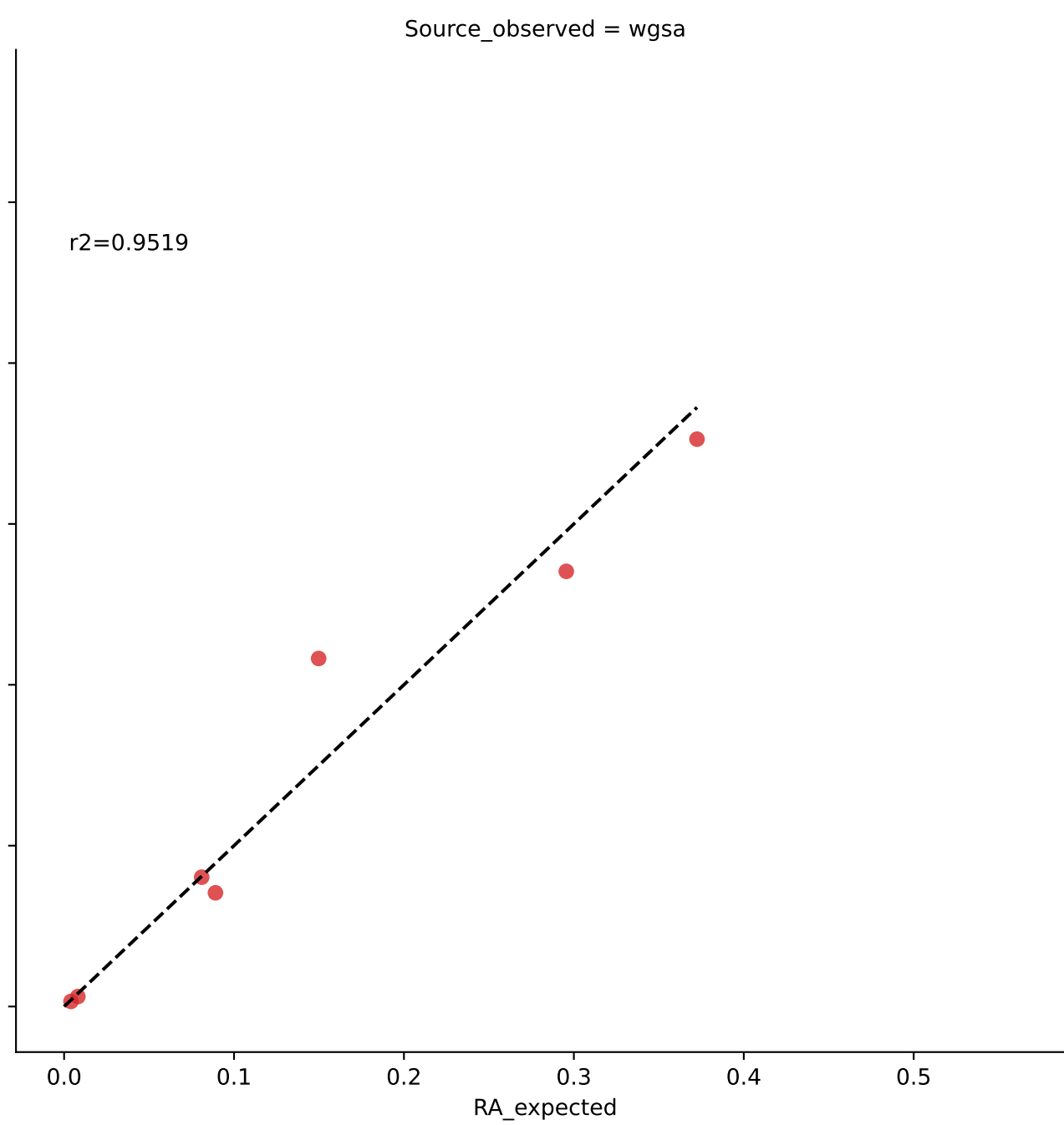
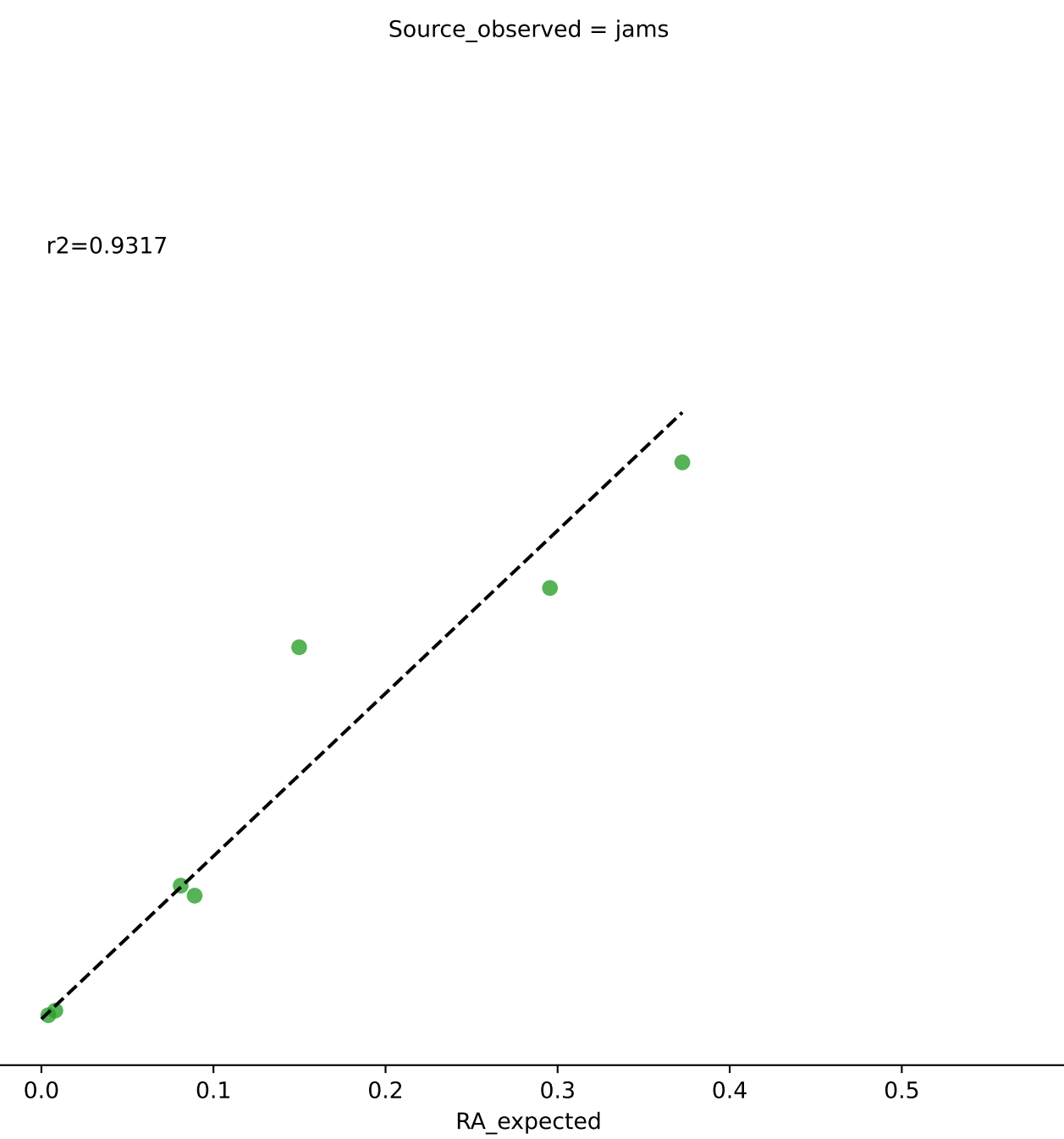
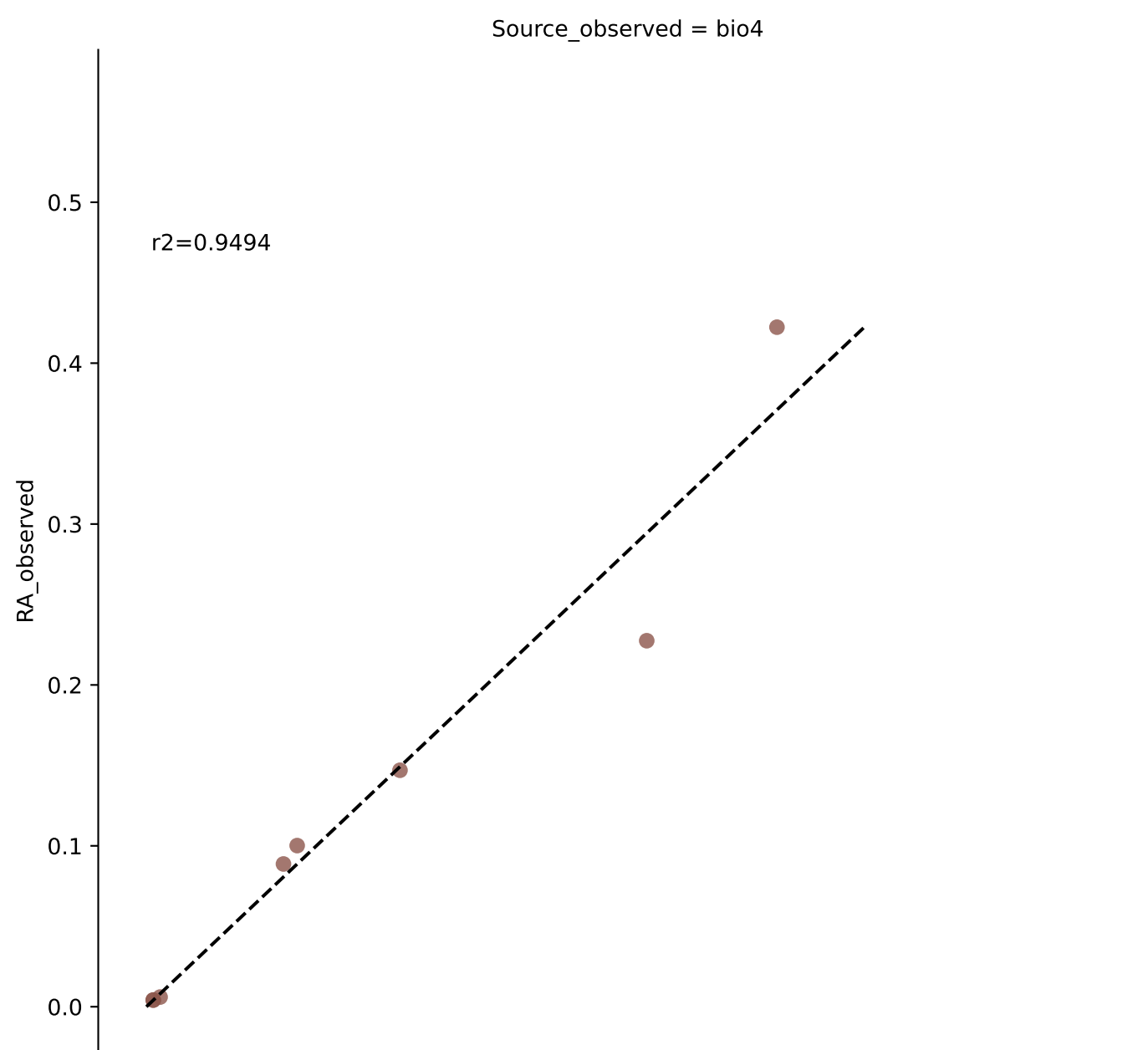


Bivariate Linear Regression for Sample S1 in Experiment camisimGI (Genus at filter threshold 0.0001)

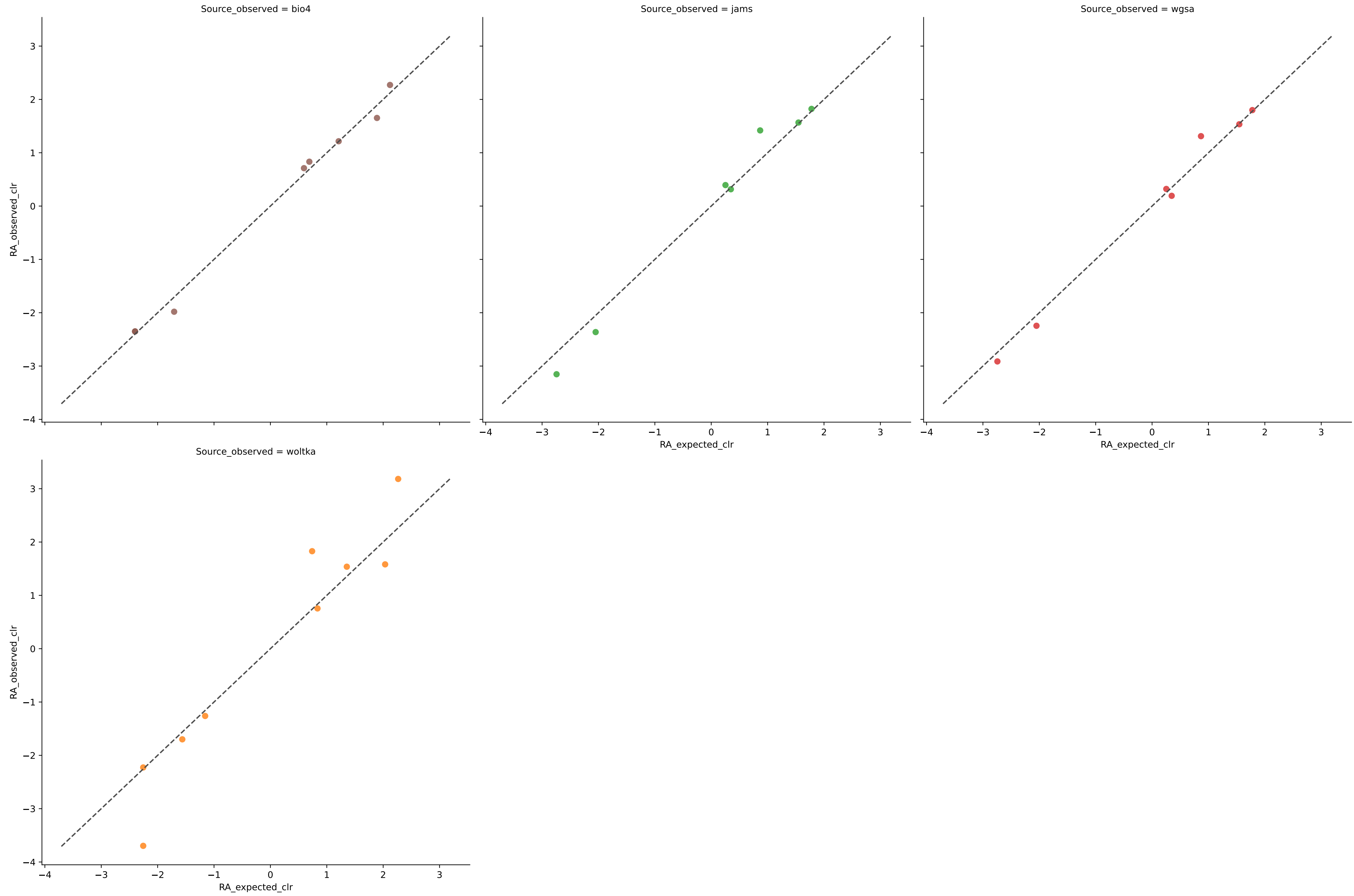


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	15	0.9954	0.0056	1.4647	0.9583	0.0124	100.0000	0.0000
jams	14	0.9995	0.0033	1.4210	0.9771	0.0047	100.0000	0.0000
wgsa	15	0.9998	0.0019	1.4092	0.9861	0.0024	100.0000	0.0000
woltka	17	0.9505	0.0143	4.1172	0.8784	0.0318	100.0000	0.0000

Bivariate Linear Regression for Sample S2 in Experiment camisimGI (Genus at filter threshold 0.0001)

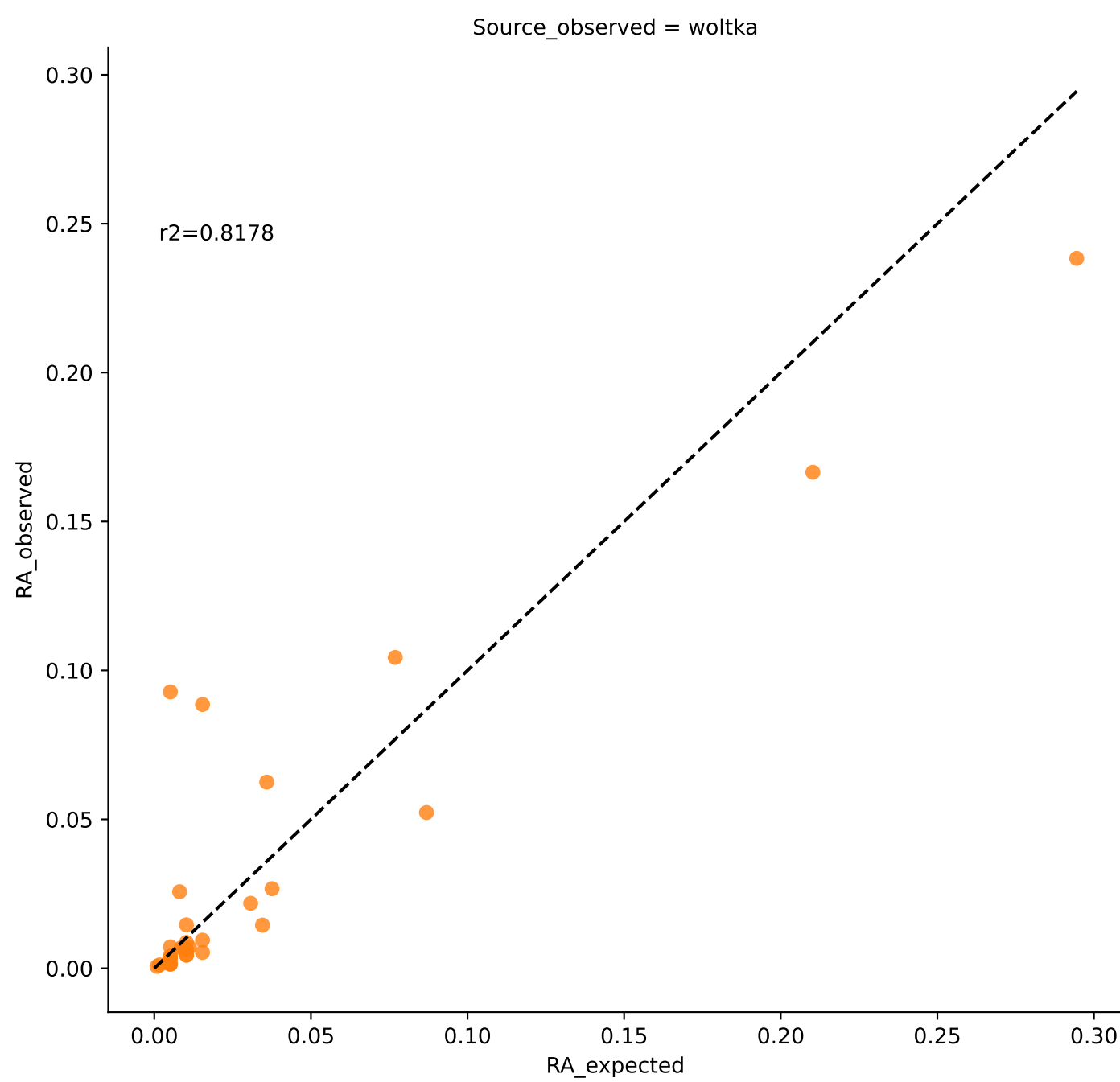
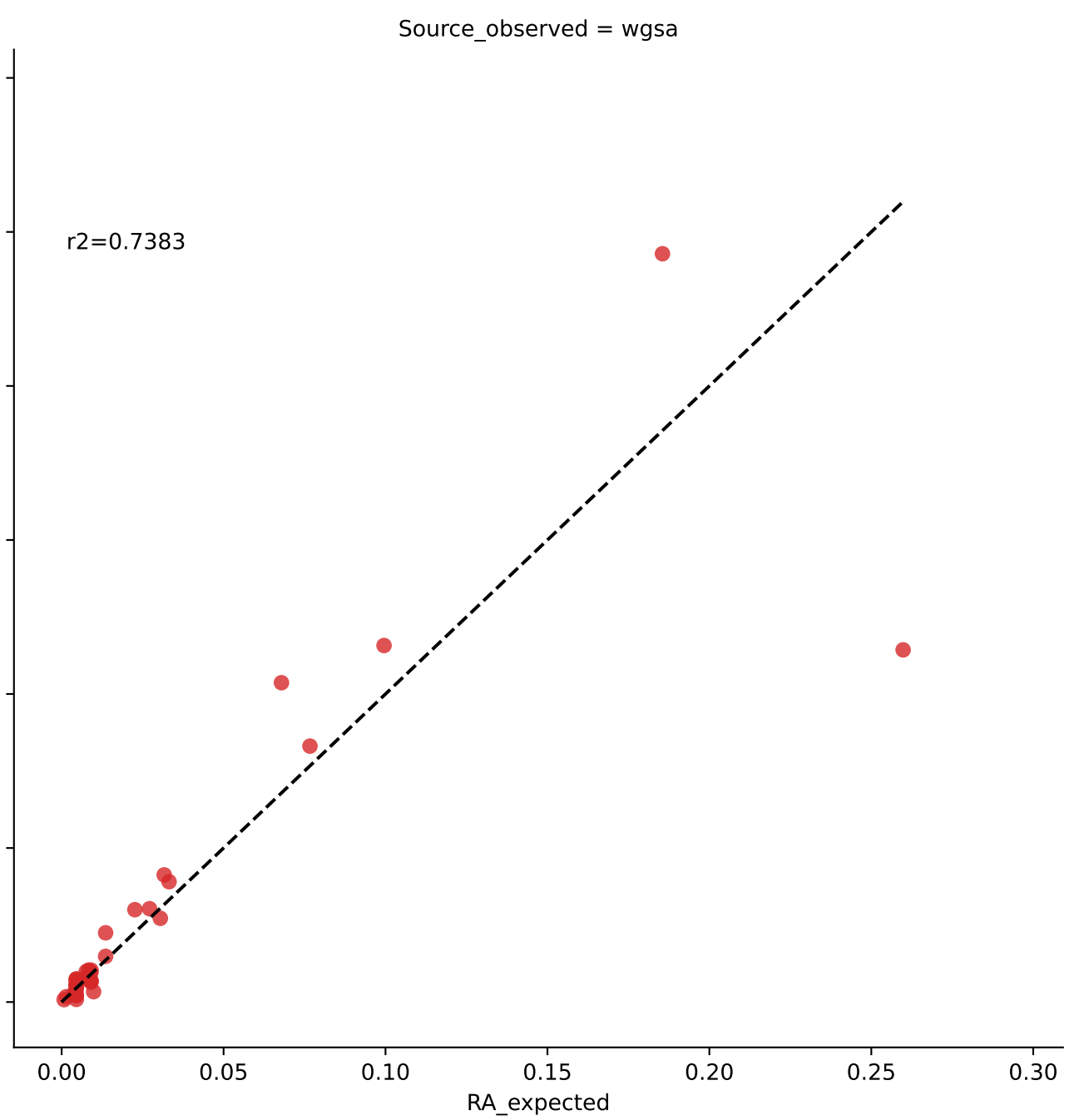
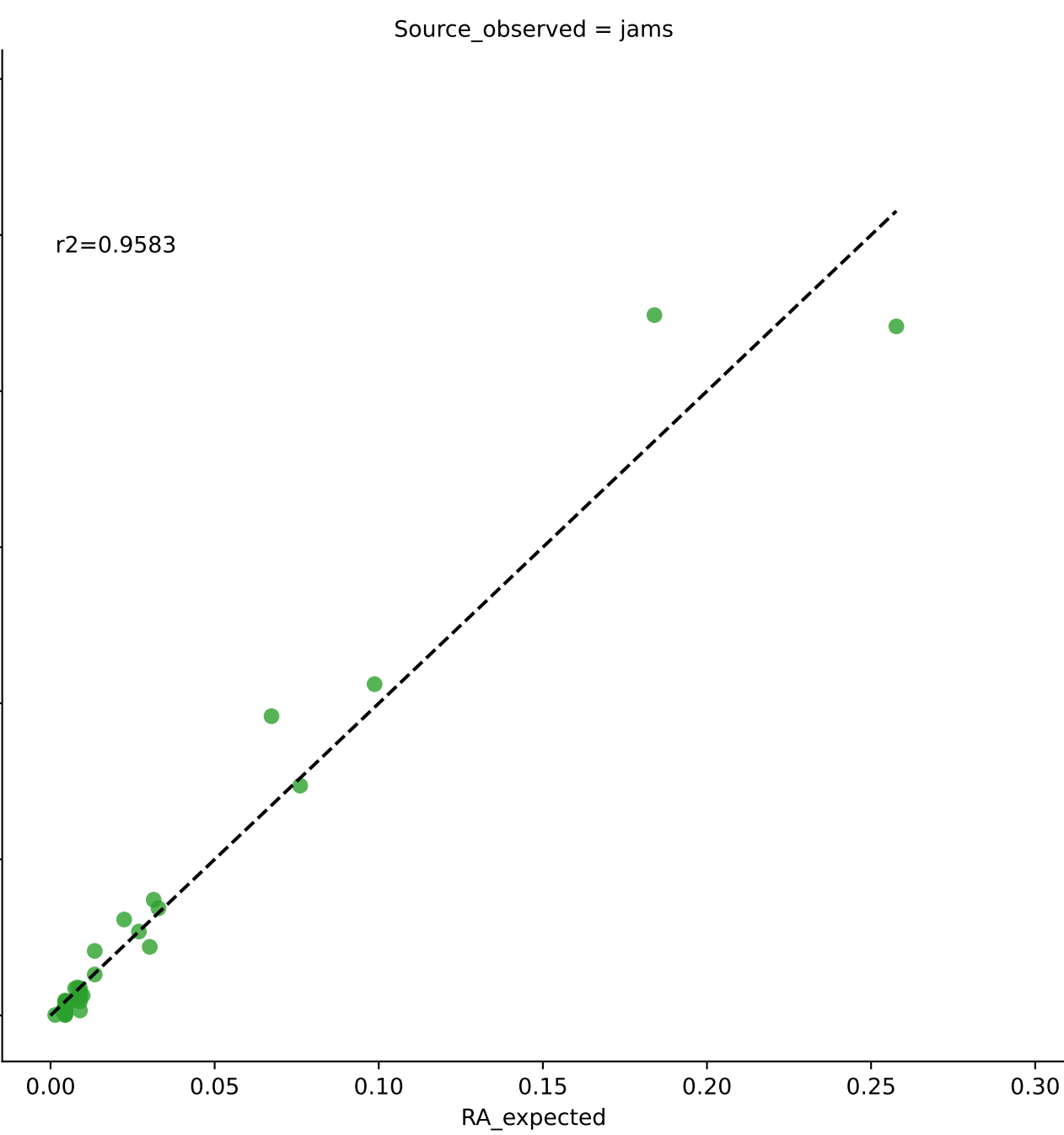
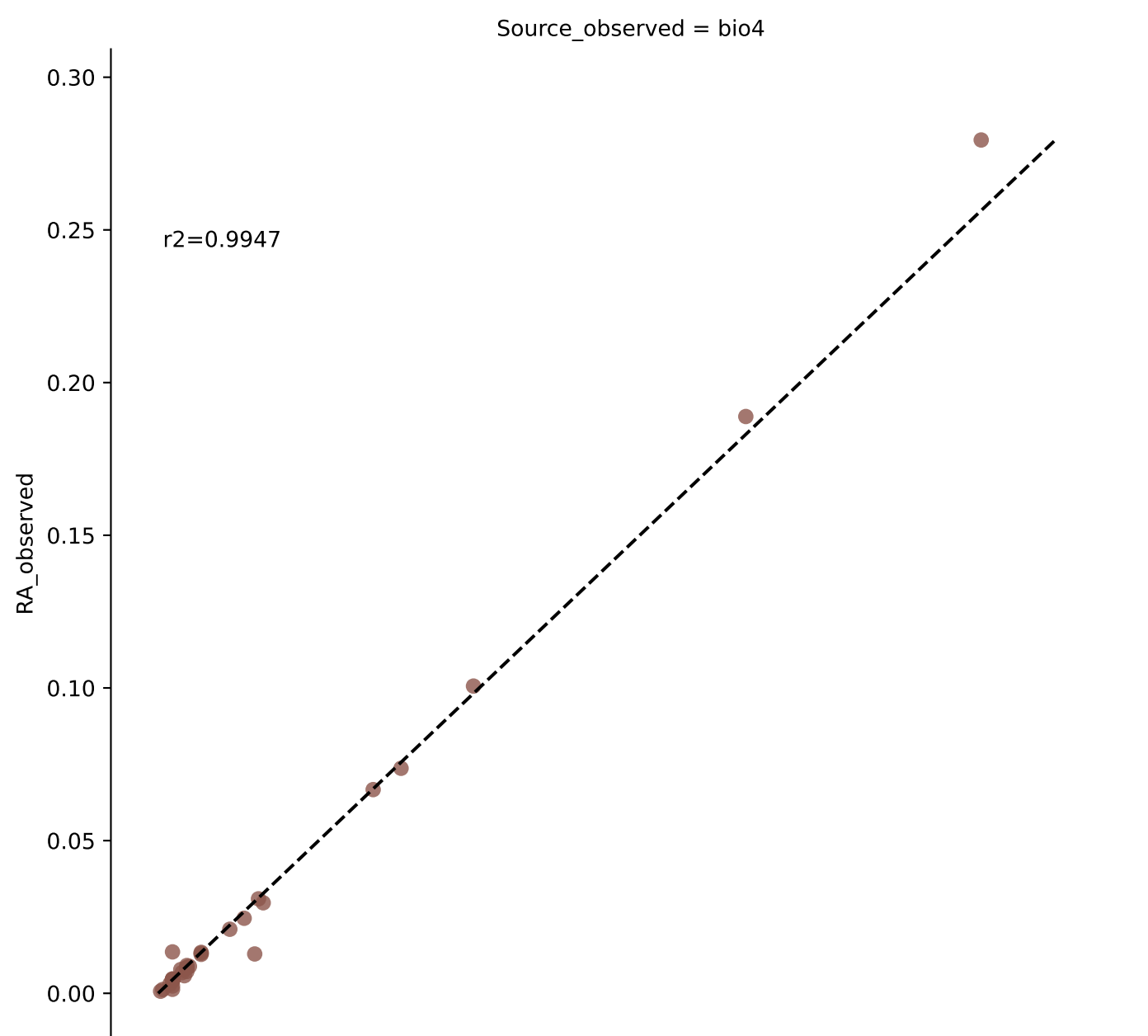


Bivariate Linear Regression for Sample S2 in Experiment camisimGI (Genus at filter threshold 0.0001)

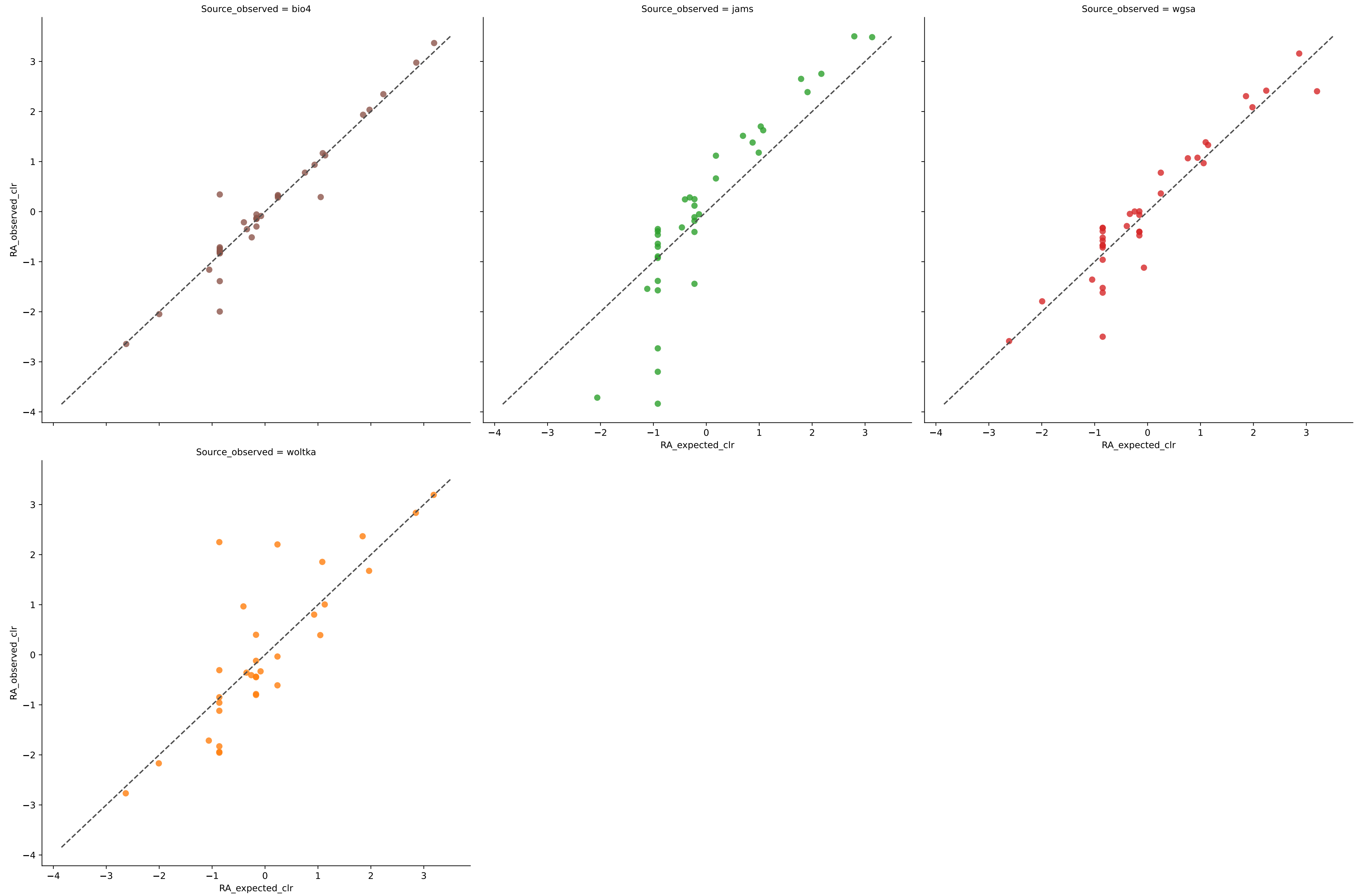


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	8	0.9494	0.0178	0.4385	0.9288	0.0302	100.0000	0.0000
jams	7	0.9317	0.0227	0.7701	0.9206	0.0343	100.0000	0.0000
wgsa	7	0.9519	0.0190	0.5411	0.9334	0.0288	100.0000	0.0000
woltka	9	0.7052	0.0592	2.0925	0.7336	0.0934	100.0000	0.0000

Bivariate Linear Regression for Sample S1 in Experiment camisimGI (Species at filter threshold 0.0001)

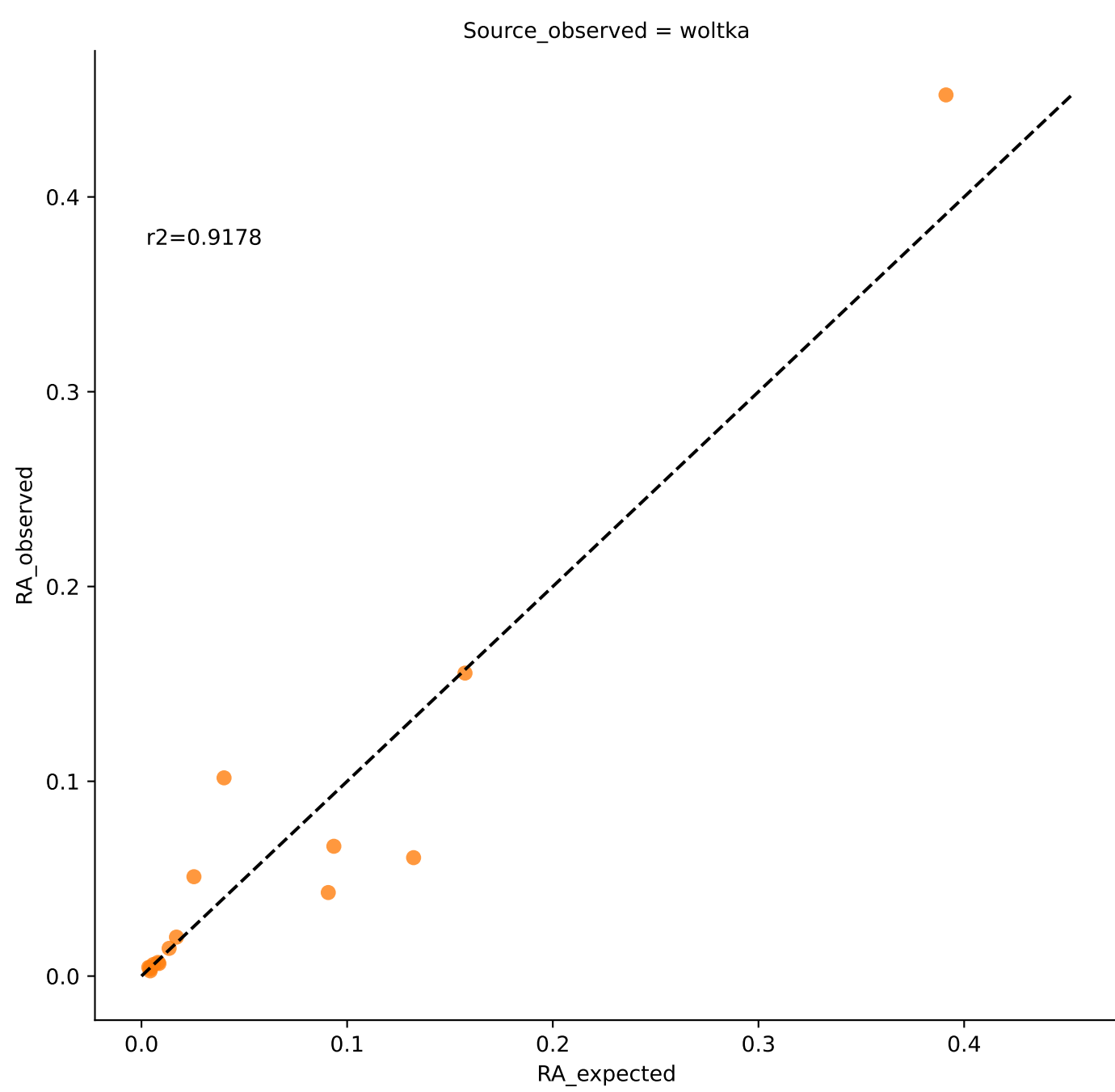
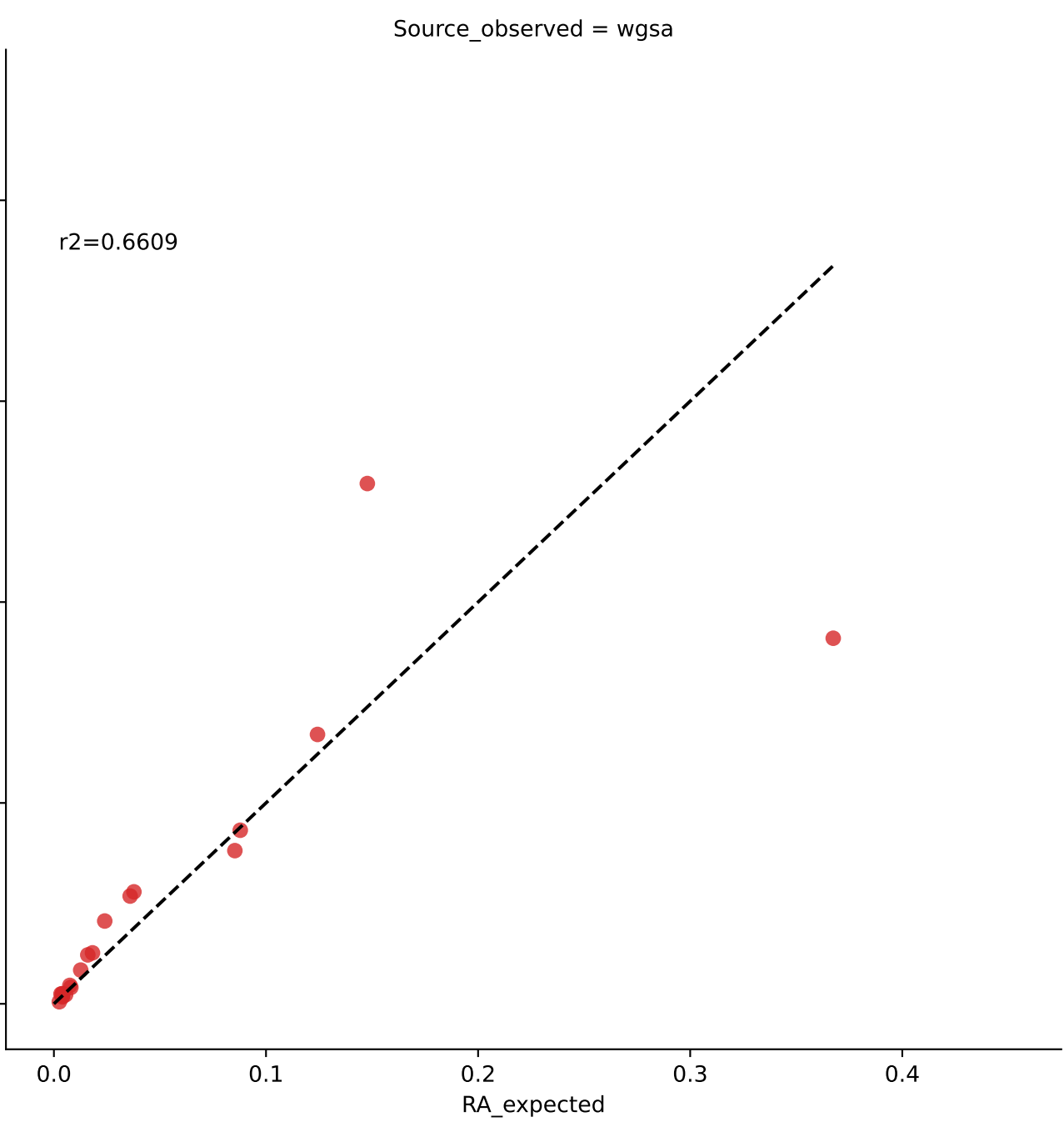
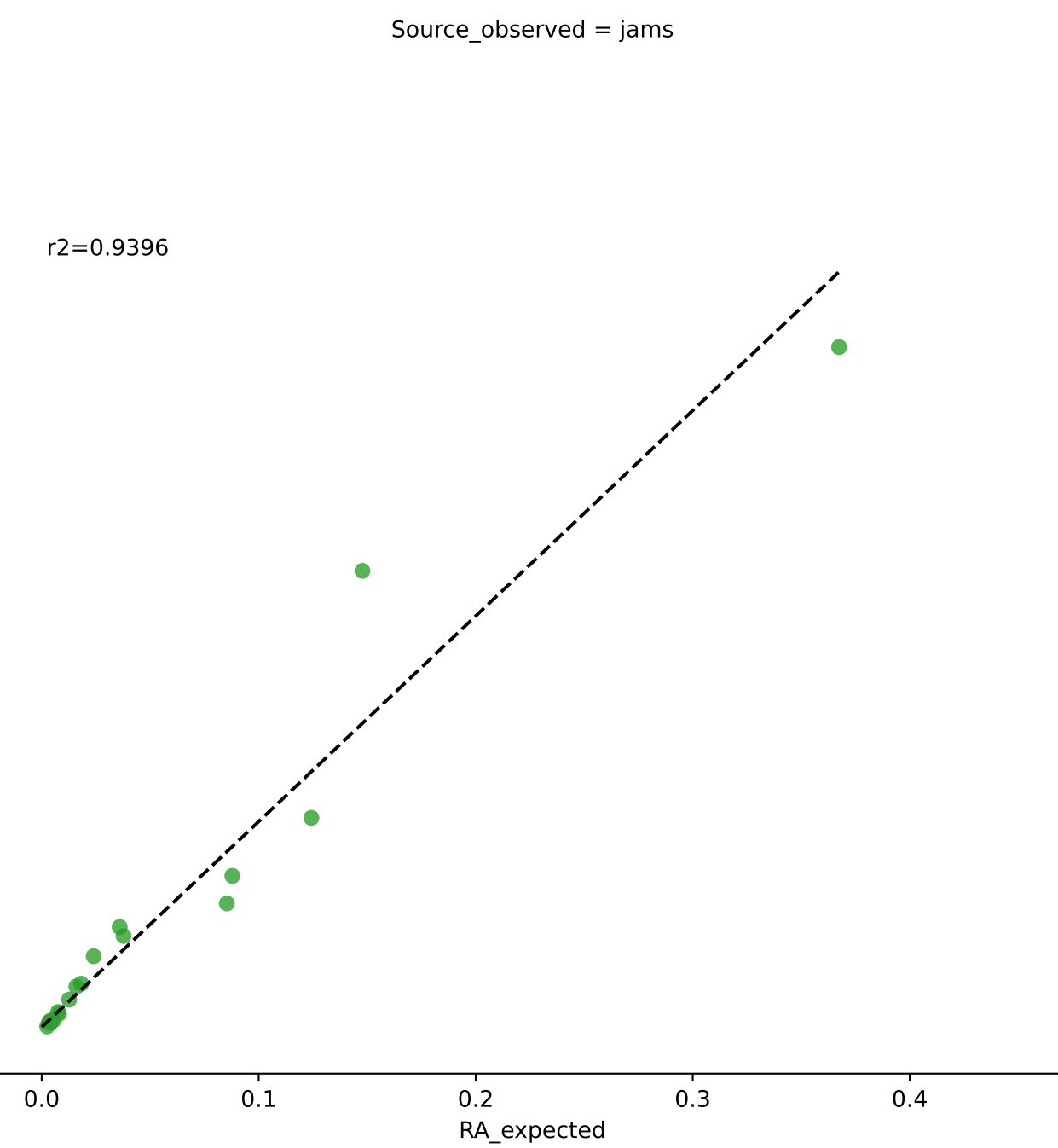
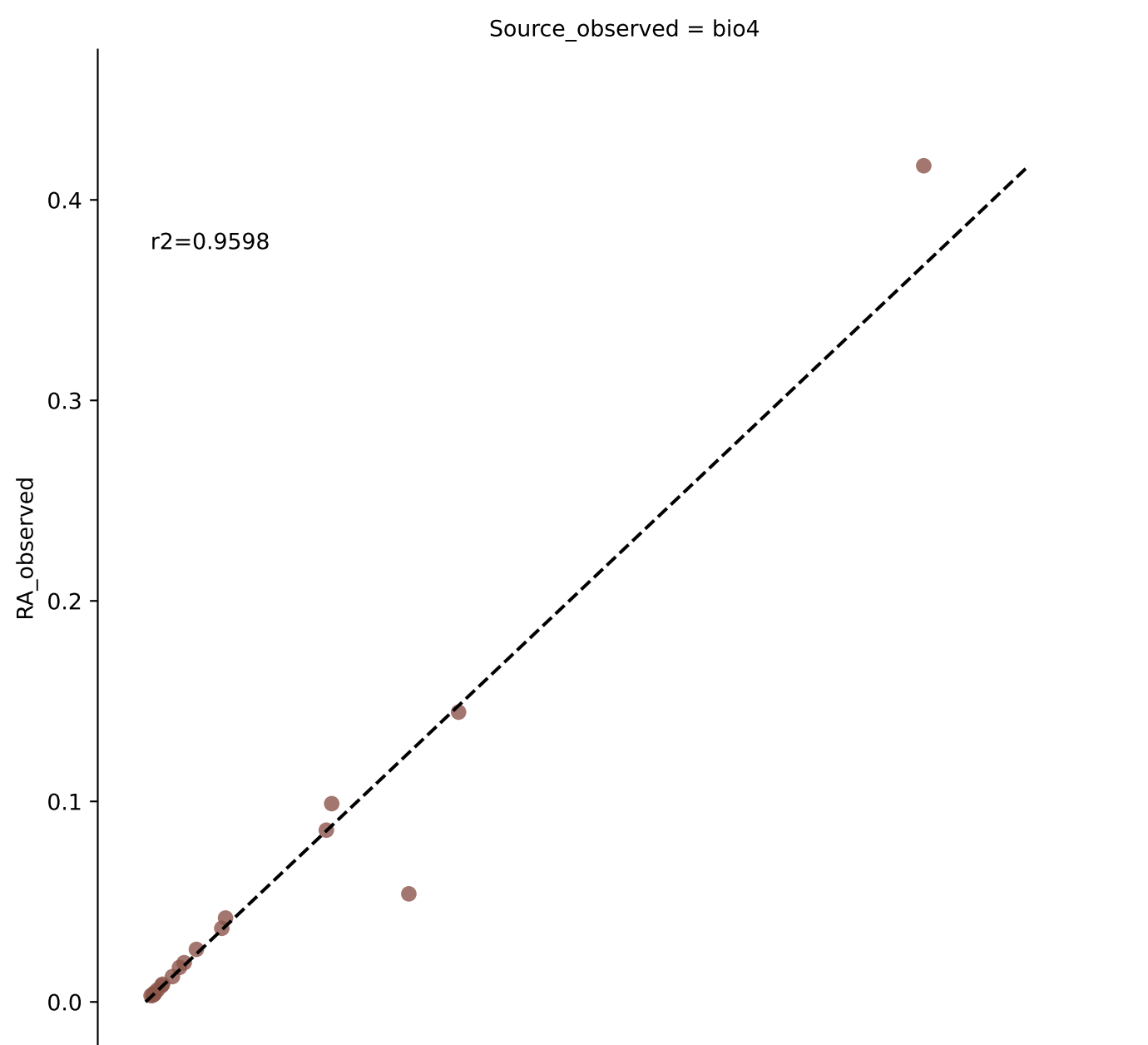


Bivariate Linear Regression for Sample S1 in Experiment camisimGI (Species at filter threshold 0.0001)

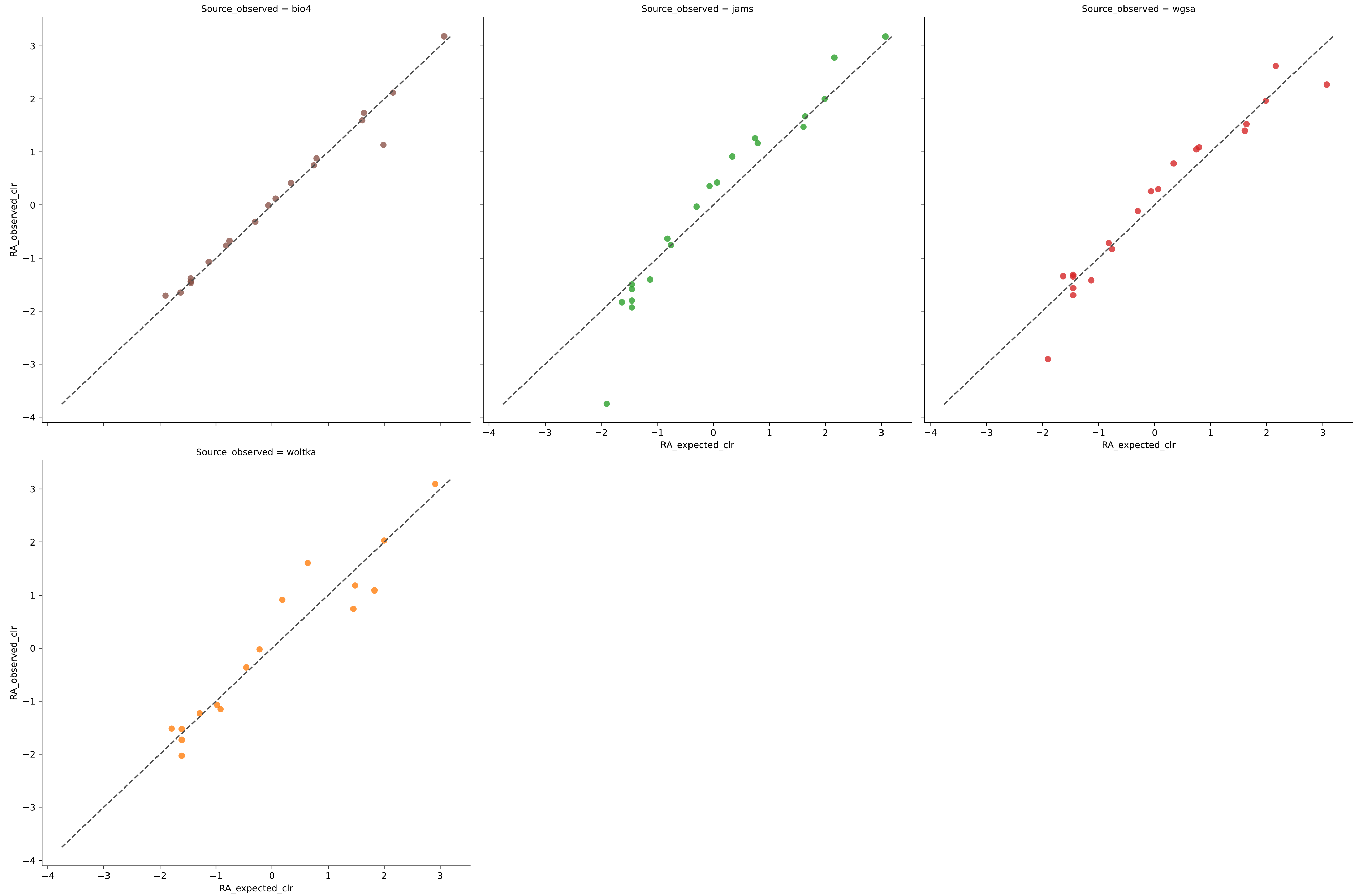


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	37	0.9947	0.0023	1.9735	0.9582	0.0052	100.0000	0.0000
jams	36	0.9583	0.0056	5.3813	0.8988	0.0109	100.0000	0.0000
wgsa	36	0.7383	0.0096	2.8185	0.8277	0.0271	100.0000	0.0000
woltka	32	0.8178	0.0149	4.8097	0.7611	0.0262	100.0000	0.0000

Bivariate Linear Regression for Sample S2 in Experiment camisimGI (Species at filter threshold 0.0001)

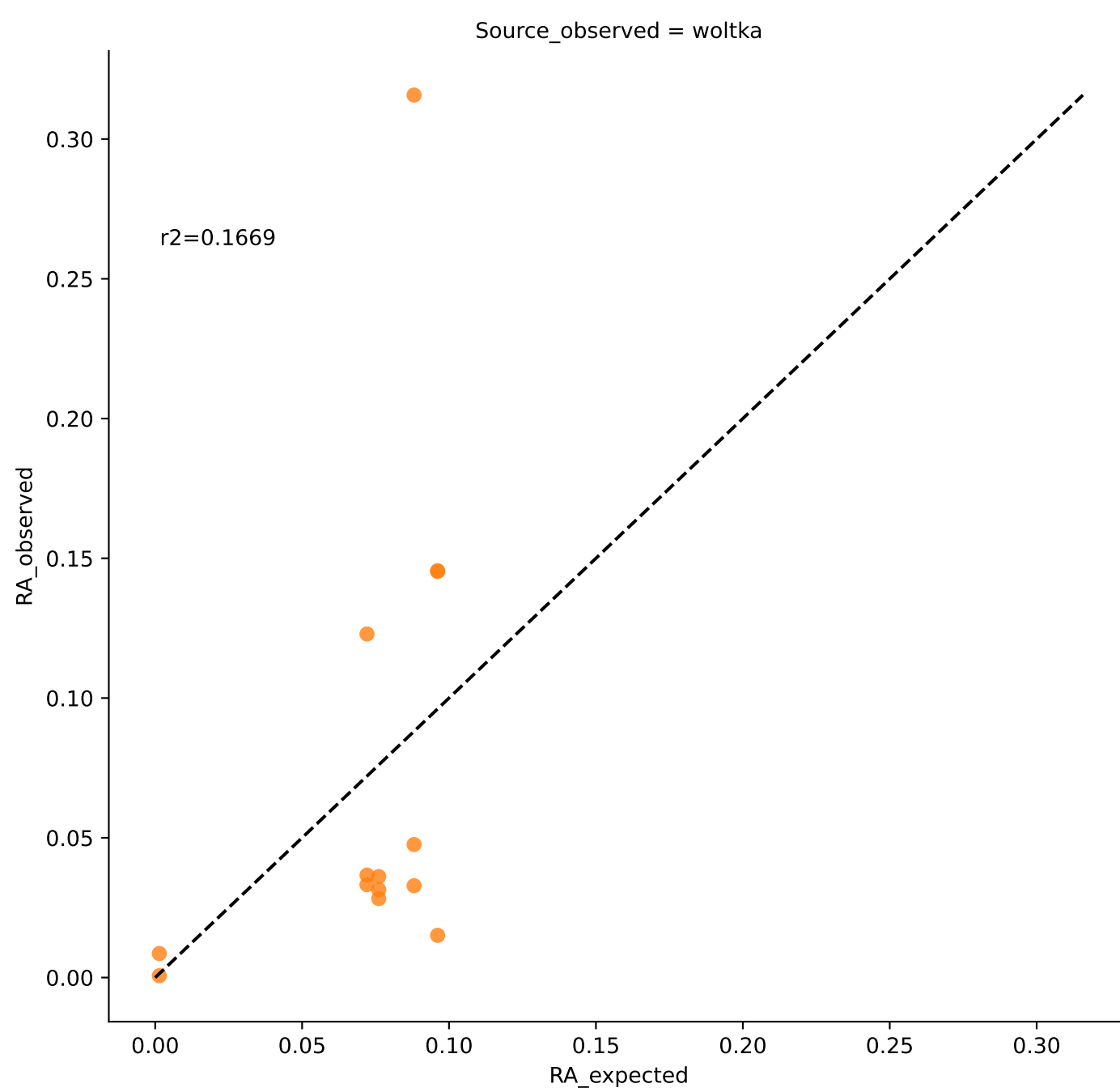
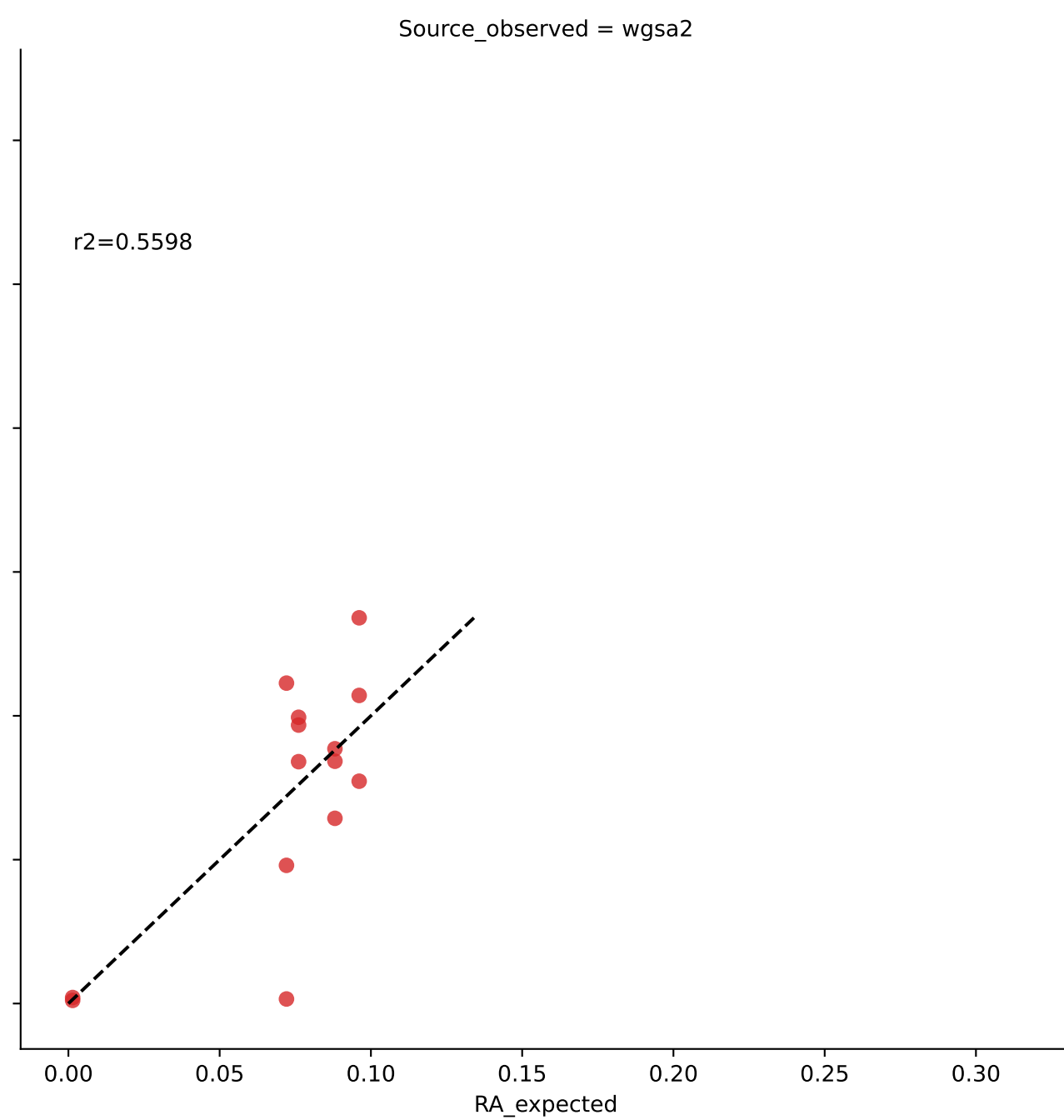
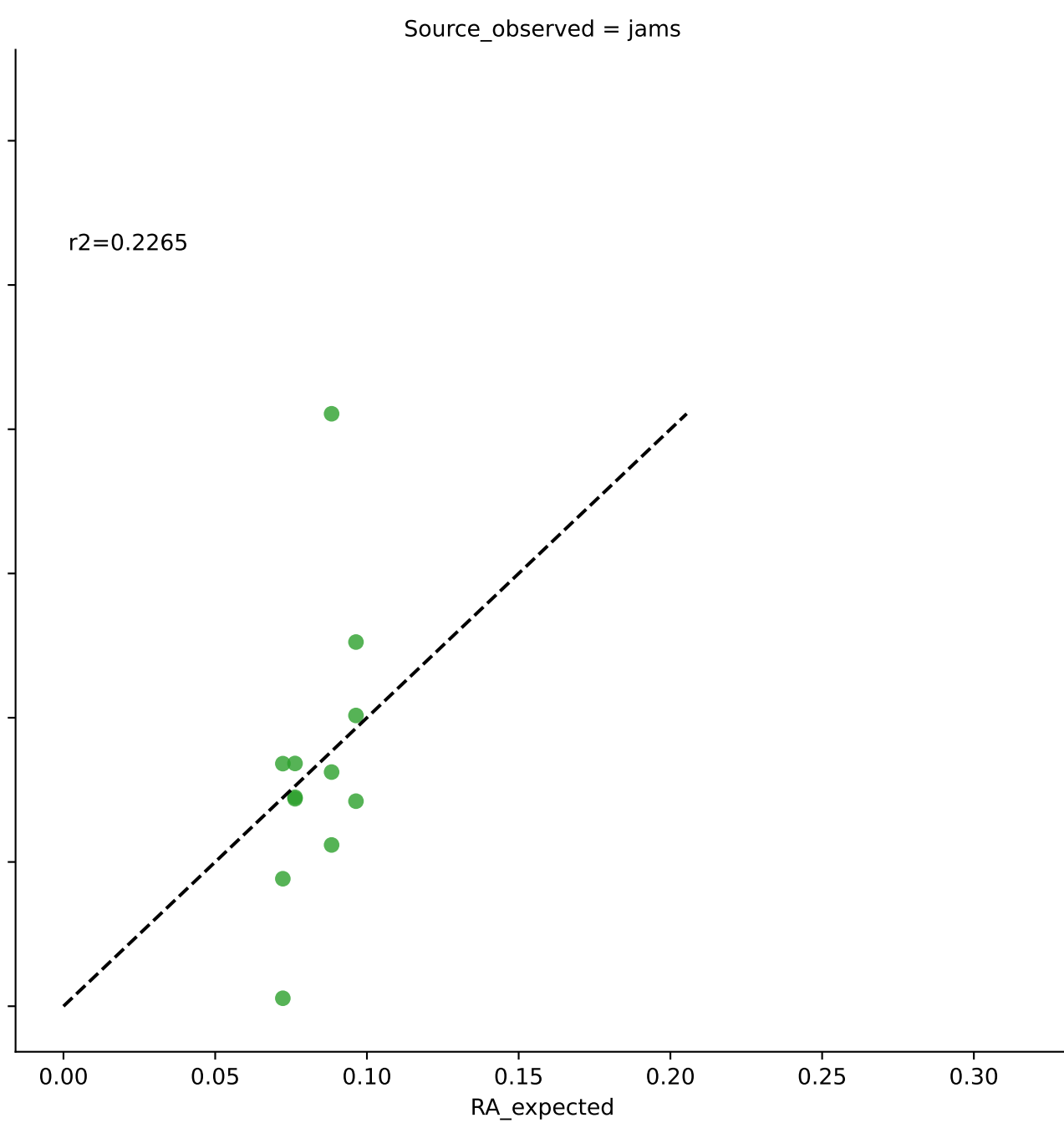
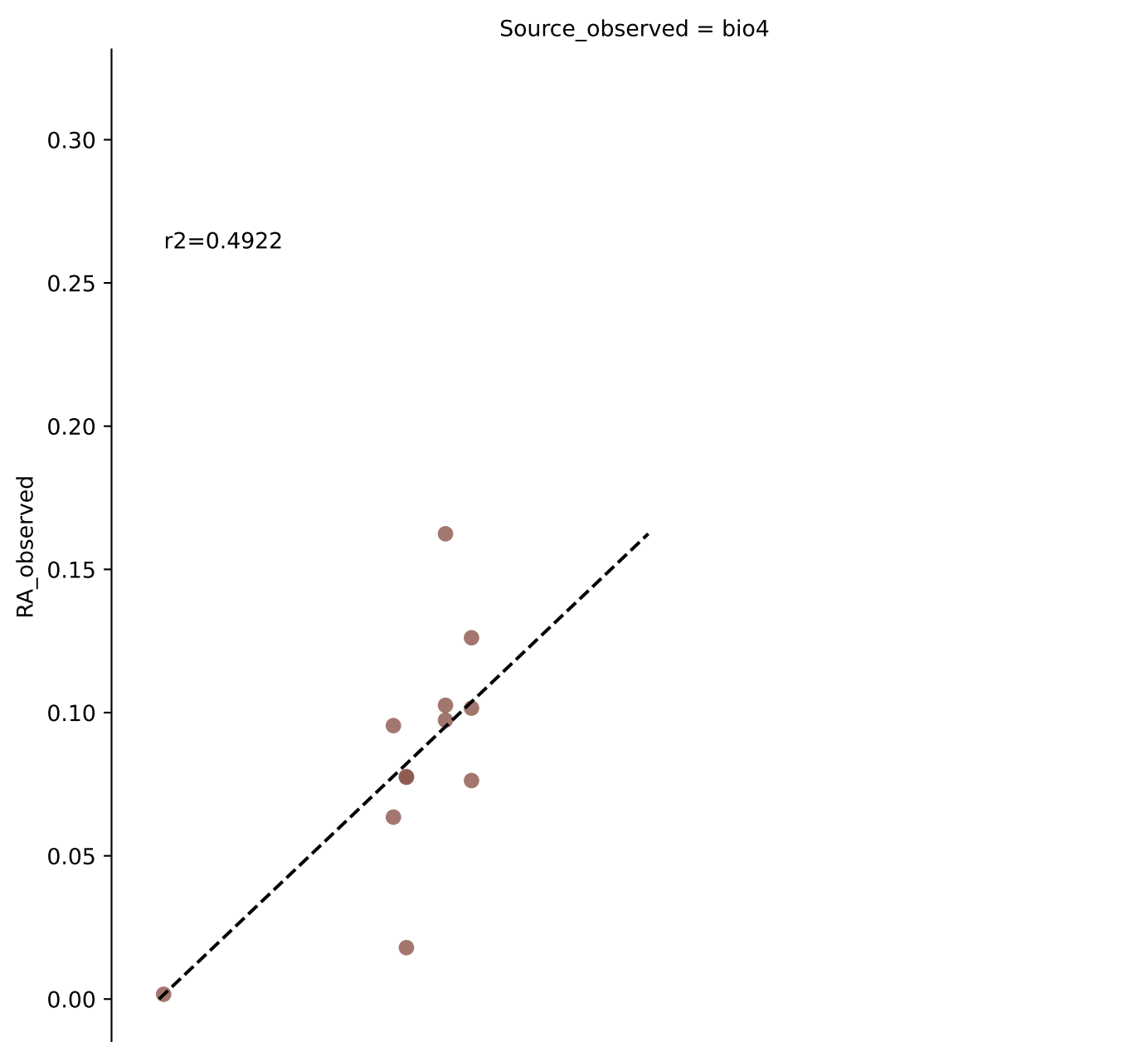


Bivariate Linear Regression for Sample S2 in Experiment camisimGI (Species at filter threshold 0.0001)

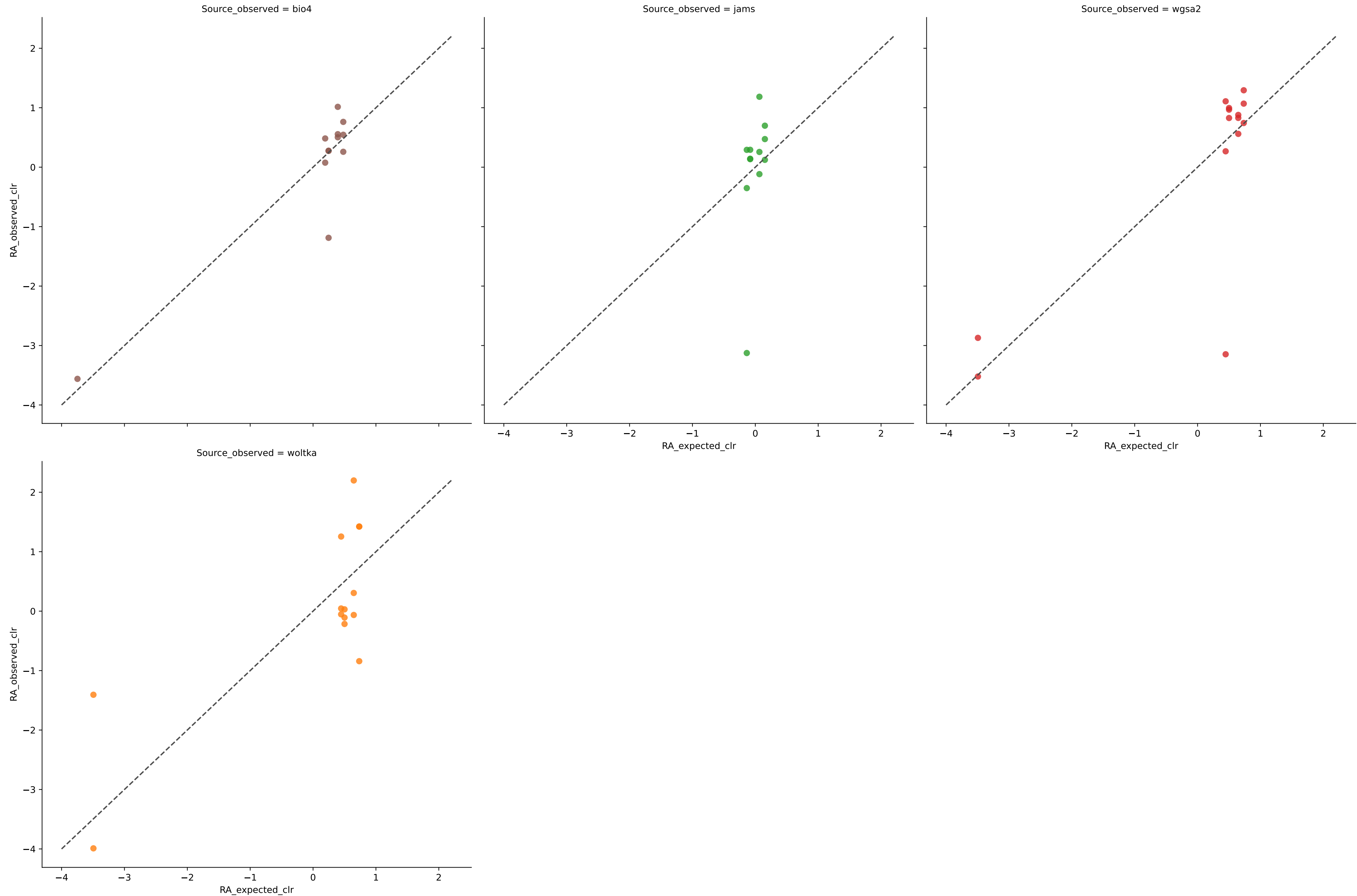


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	20	0.9598	0.0074	0.9077	0.9264	0.0195	100.0000	0.0000
jams	20	0.9396	0.0112	2.3368	0.8885	0.0207	100.0000	0.0000
wgsa	20	0.6609	0.0199	1.6681	0.8009	0.0490	100.0000	0.0000
woltka	16	0.9178	0.0192	1.7436	0.8467	0.0320	100.0000	0.0000

Bivariate Linear Regression for Sample EG in Experiment nist (Genus at filter threshold 0.0001)

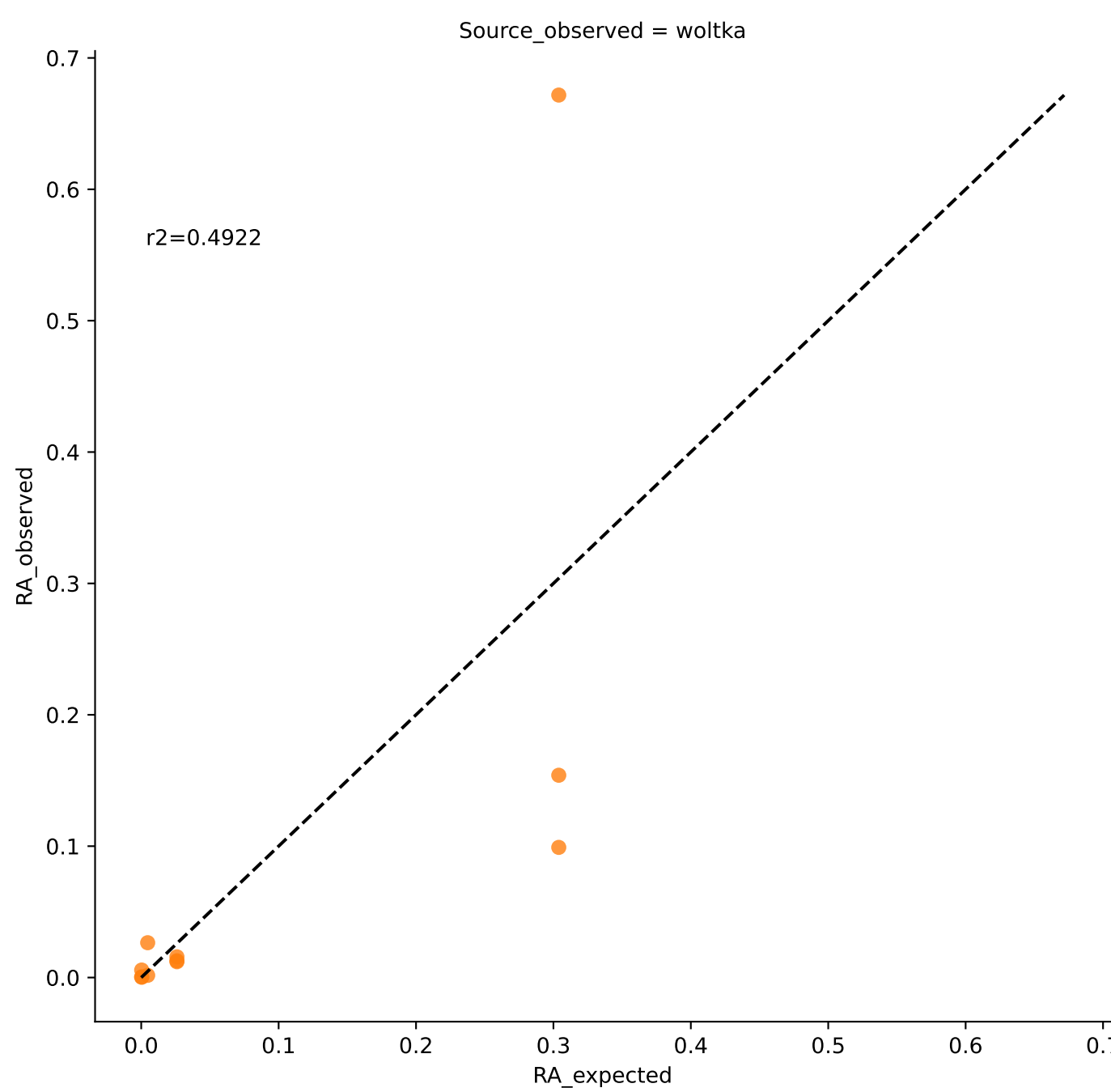
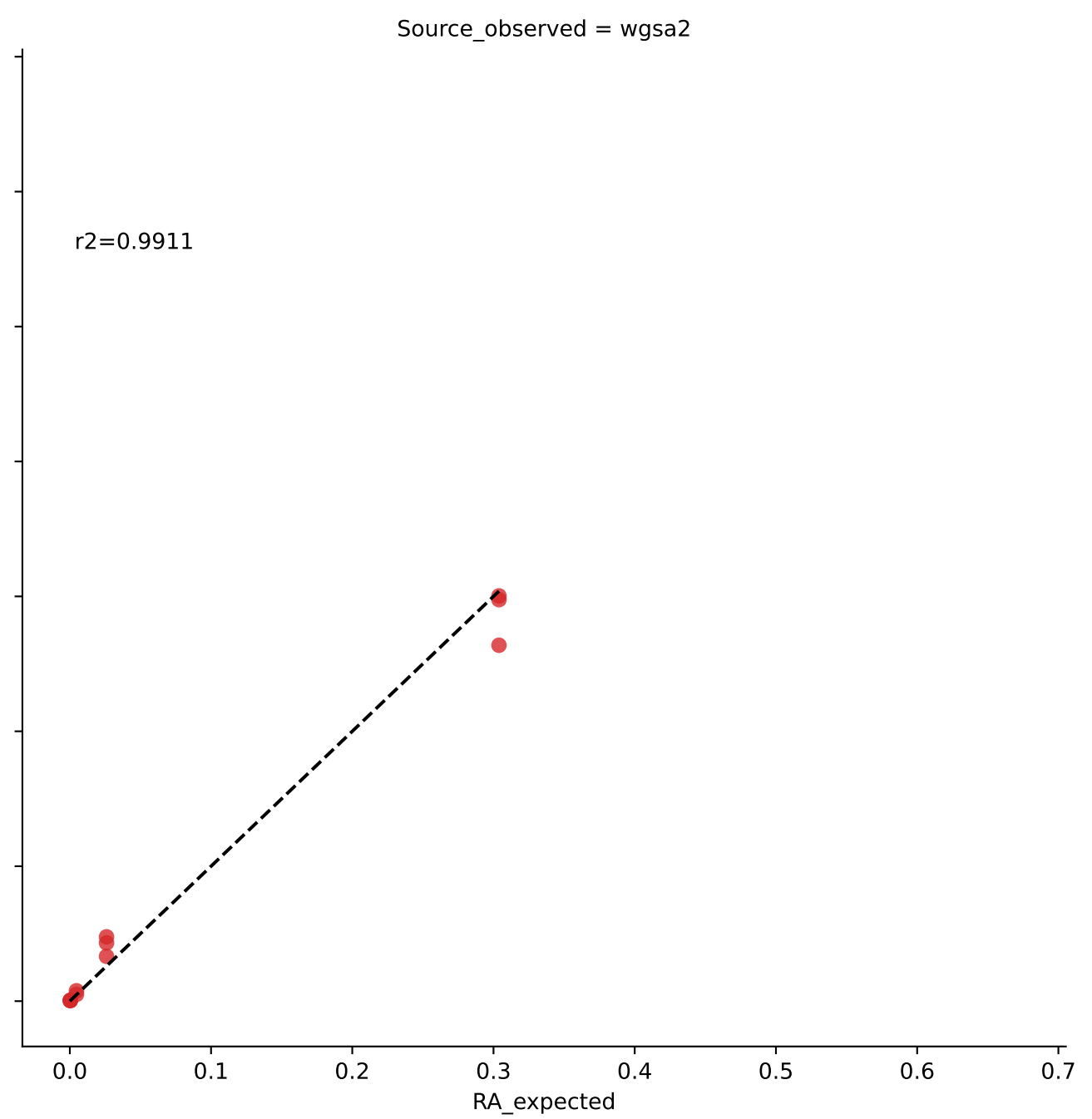
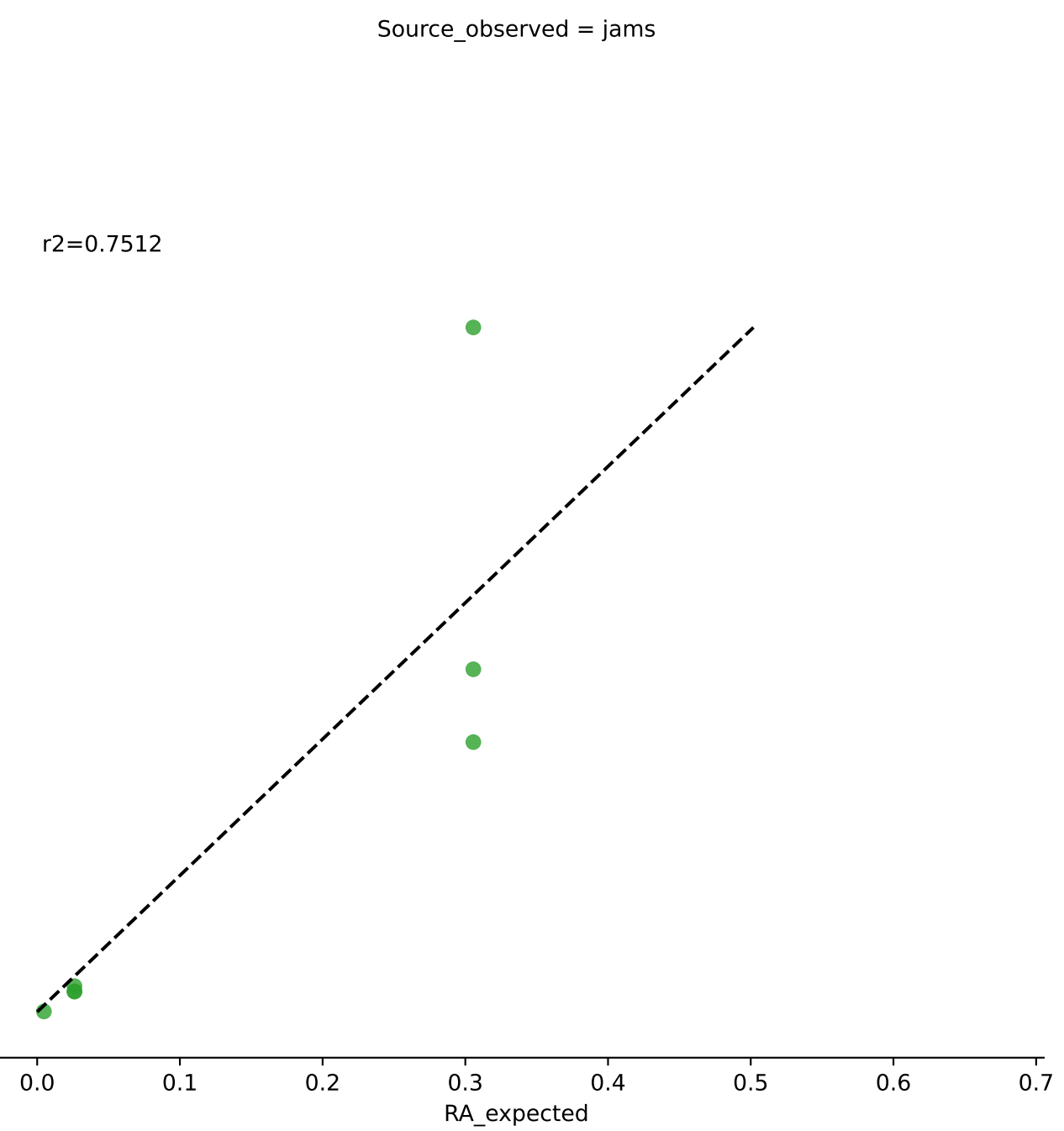
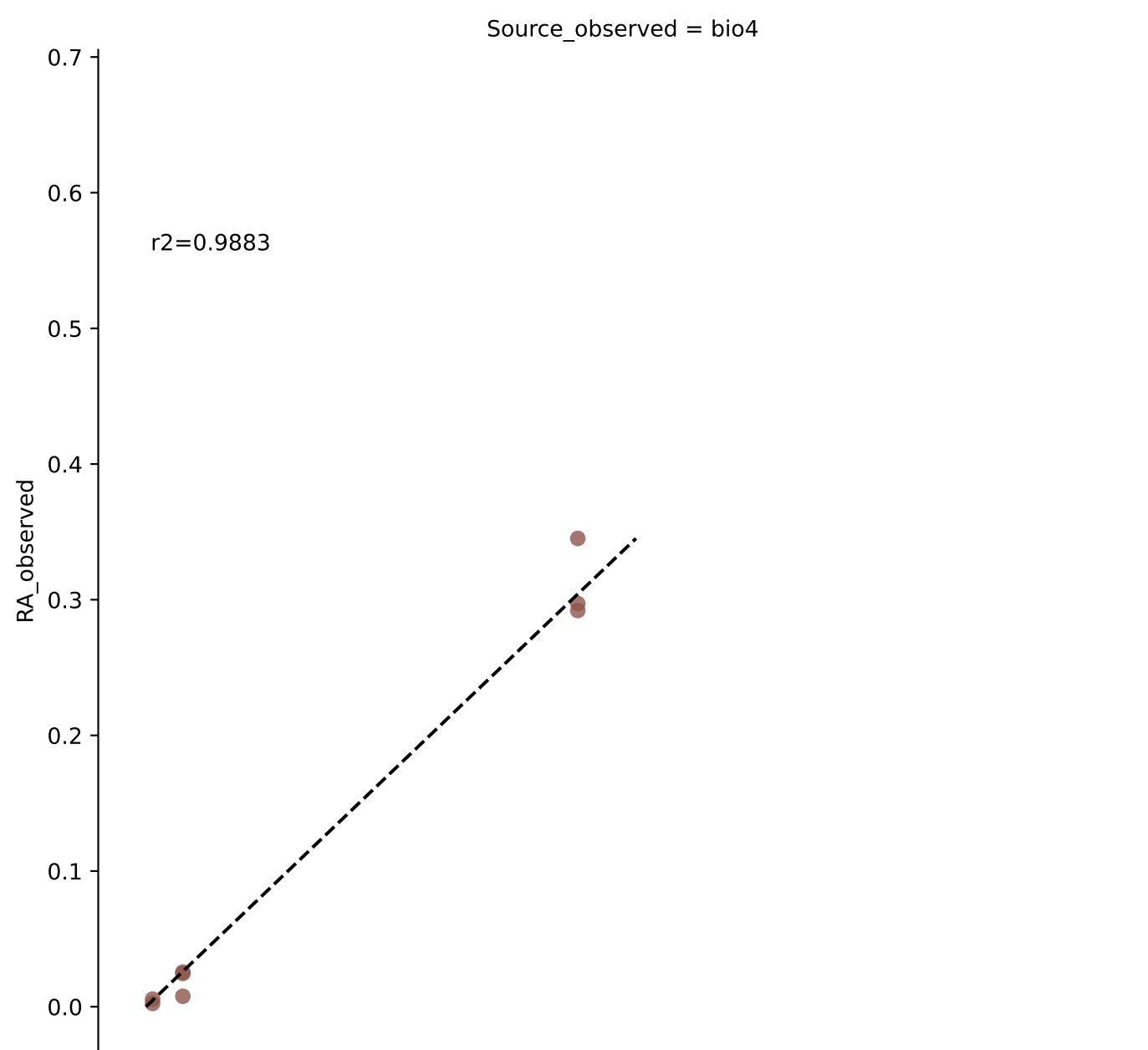


Bivariate Linear Regression for Sample EG in Experiment nist (Genus at filter threshold 0.0001)

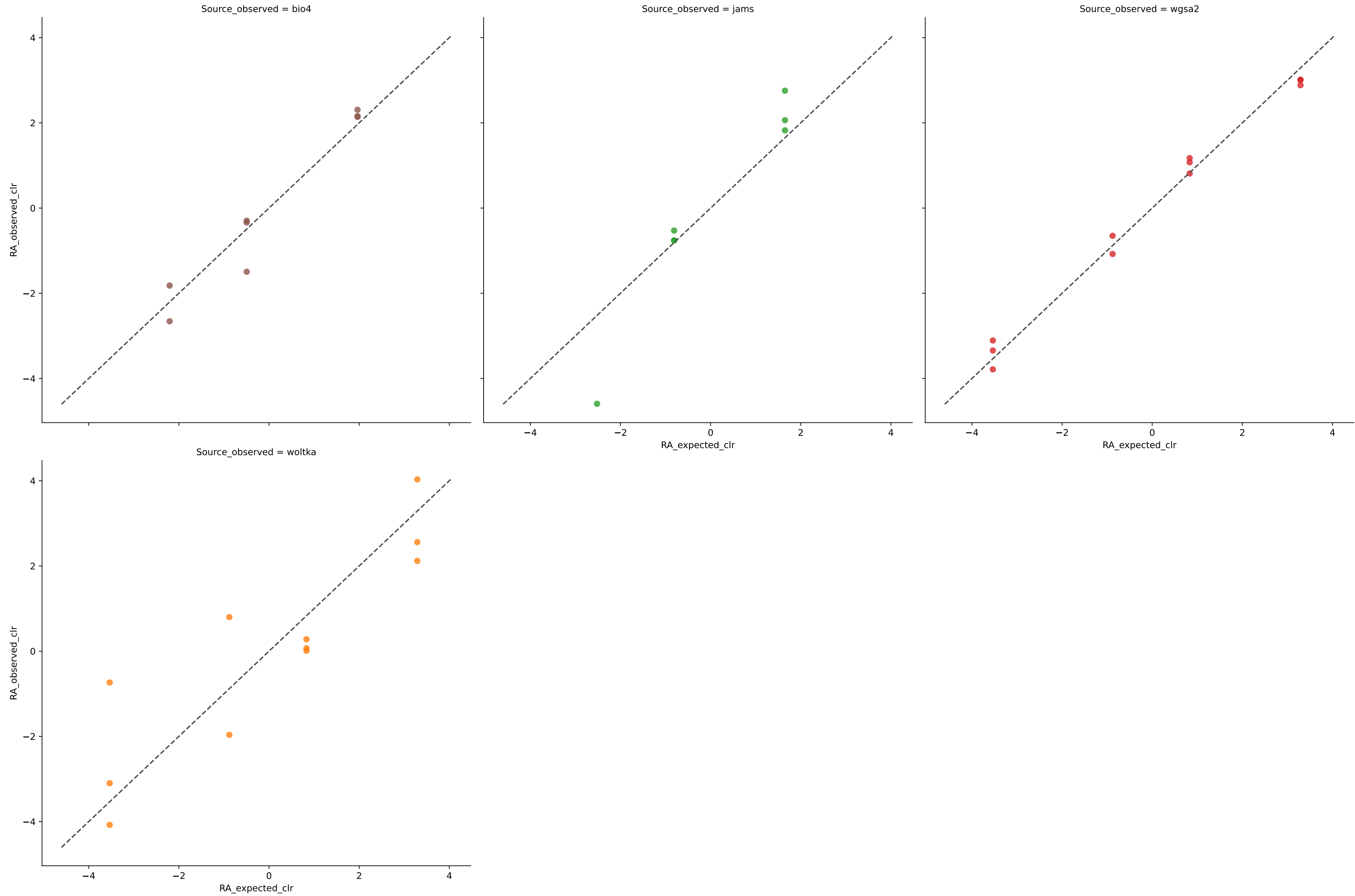


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	12	0.4922	0.0195	1.6584	0.8827	0.0296	100.0000	0.0000
jams	12	0.2265	0.0285	3.3366	0.8291	0.0430	100.0000	0.0000
wgsa2	14	0.5598	0.0202	3.8550	0.8586	0.0276	100.0000	0.0000
woltka	14	0.1669	0.0549	3.6377	0.6157	0.0752	100.0000	0.0000

Bivariate Linear Regression for Sample MIX-A in Experiment nist (Genus at filter threshold 0.0001)

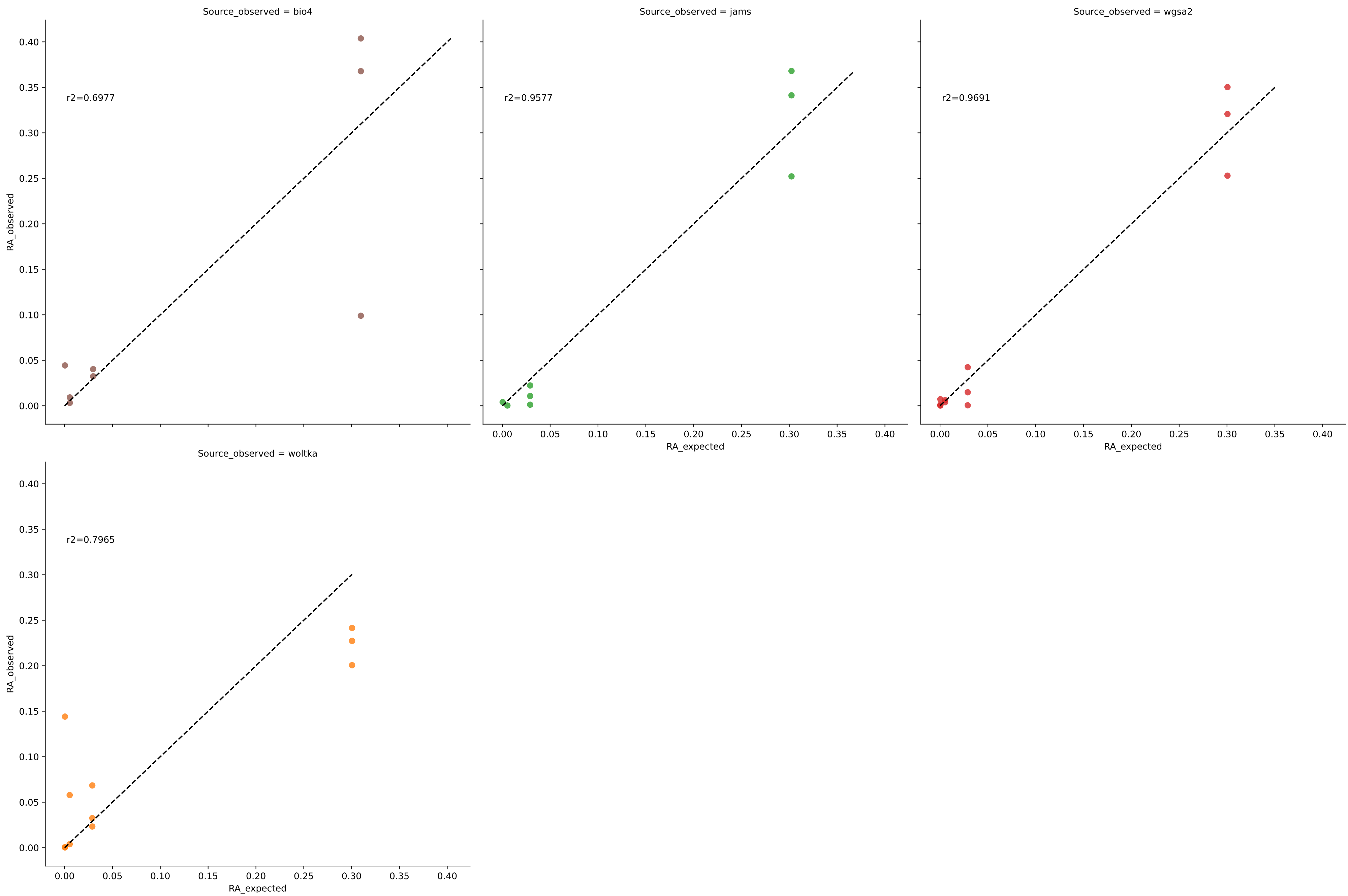


Bivariate Linear Regression for Sample MIX-A in Experiment nist (Genus at filter threshold 0.0001)

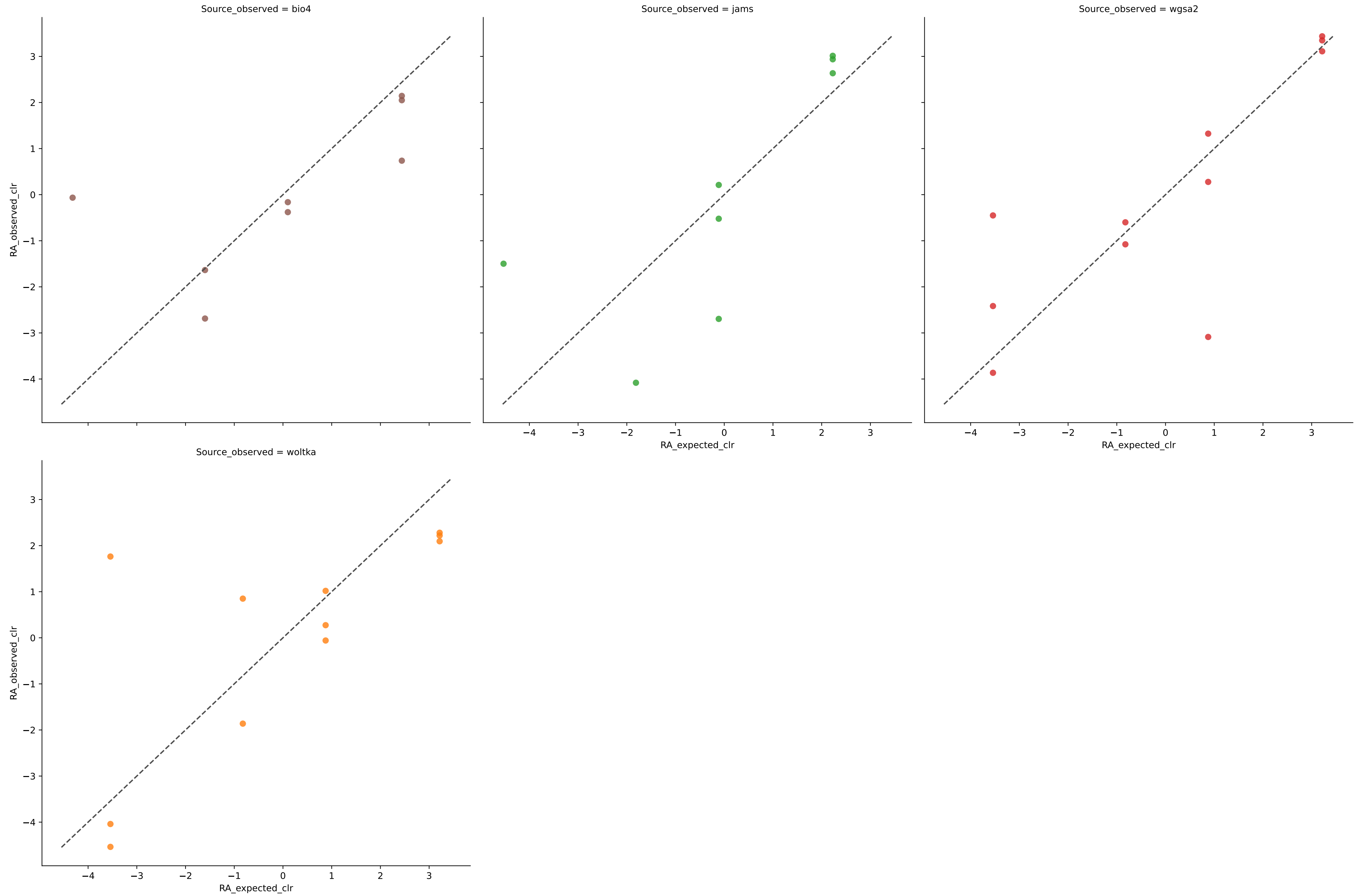


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	8	0.9883	0.0105	1.2667	0.9582	0.0167	100.0000	0.0000
jams	7	0.7512	0.0561	2.4101	0.8037	0.0874	100.0000	0.0000
wgsa2	11	0.9911	0.0091	0.9344	0.9501	0.0150	100.0000	0.0000
woltka	11	0.4922	0.0719	4.0410	0.6048	0.1351	100.0000	0.0000

Bivariate Linear Regression for Sample MIX-B in Experiment nist (Genus at filter threshold 0.0001)

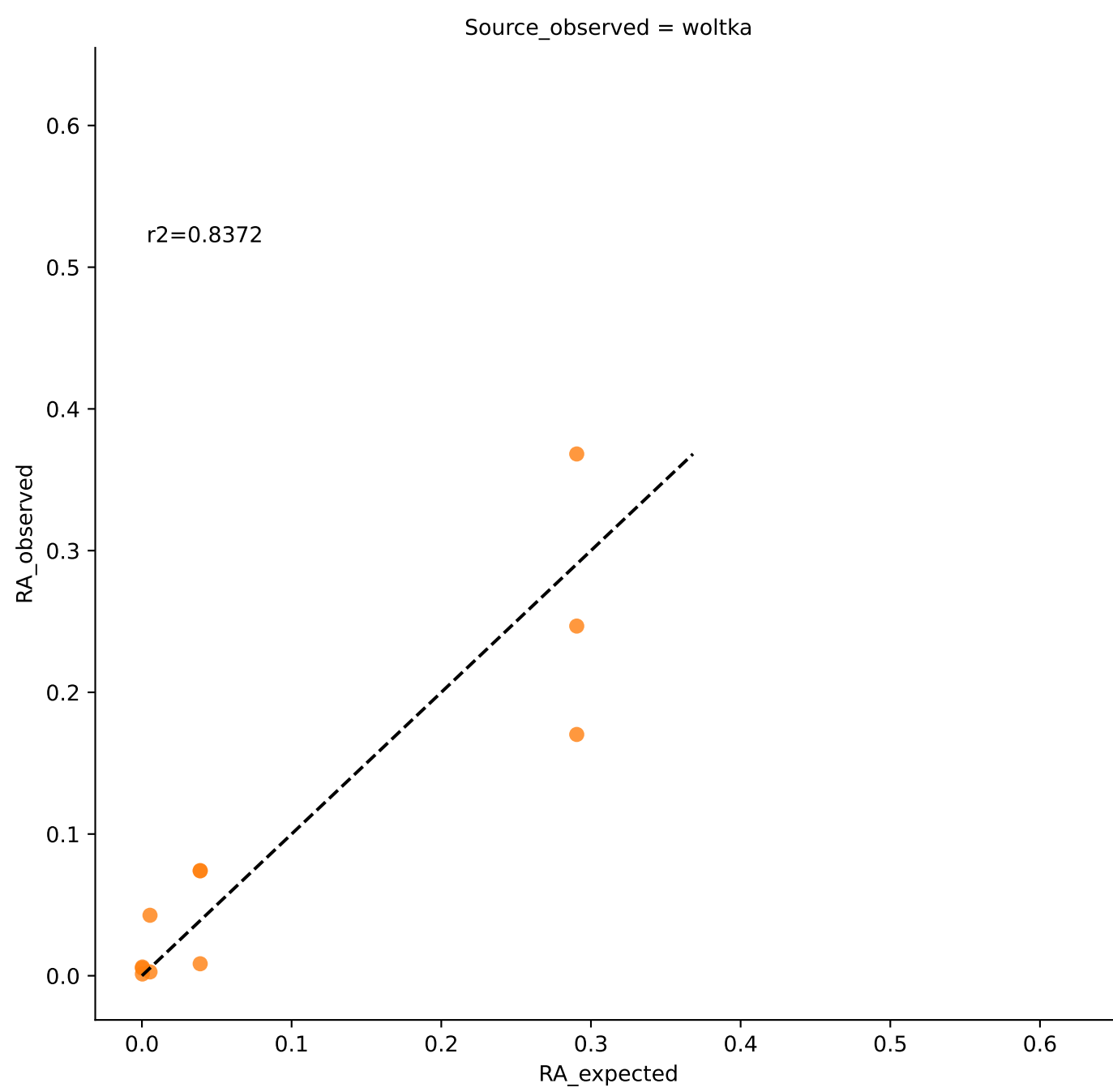
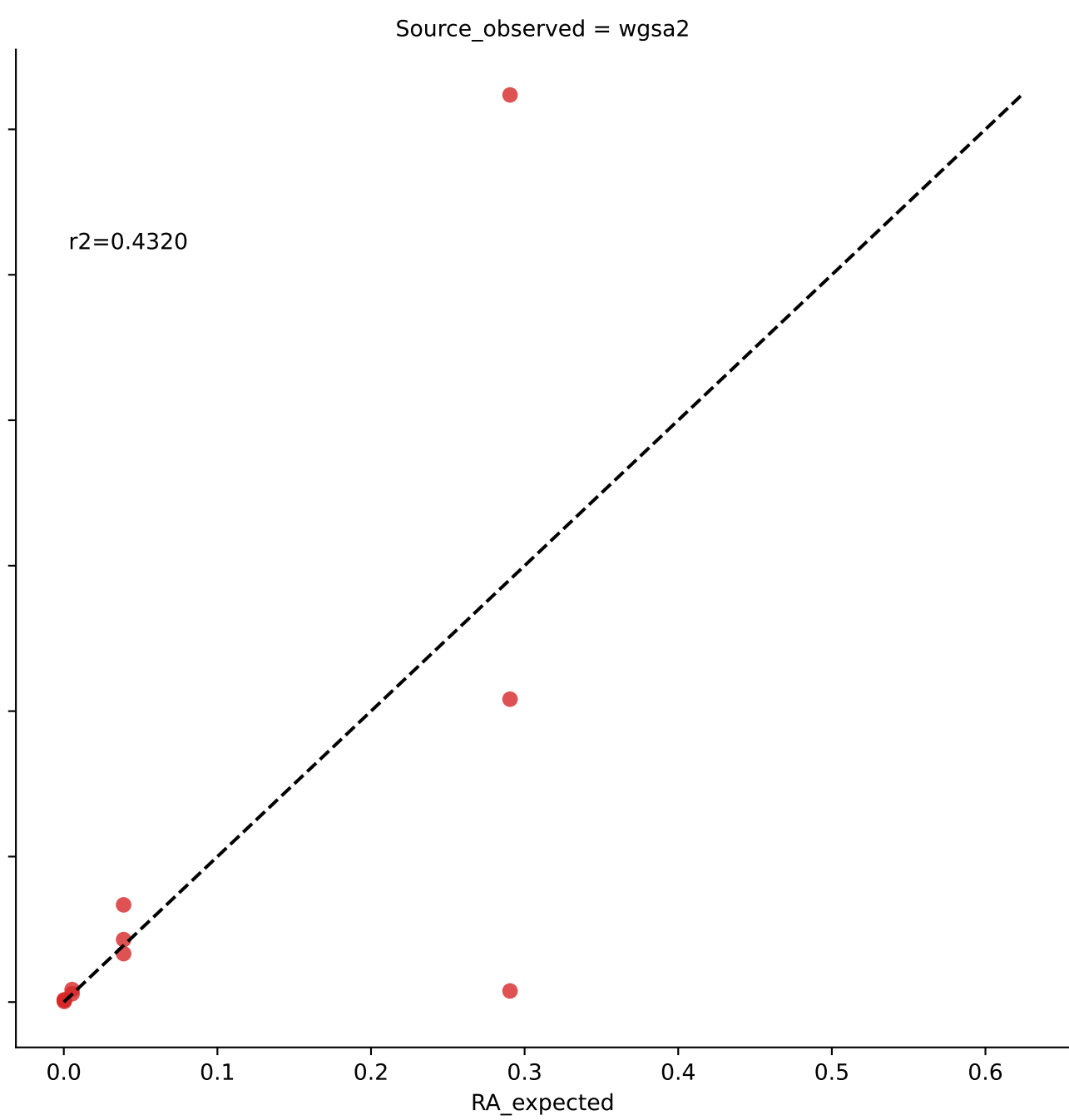
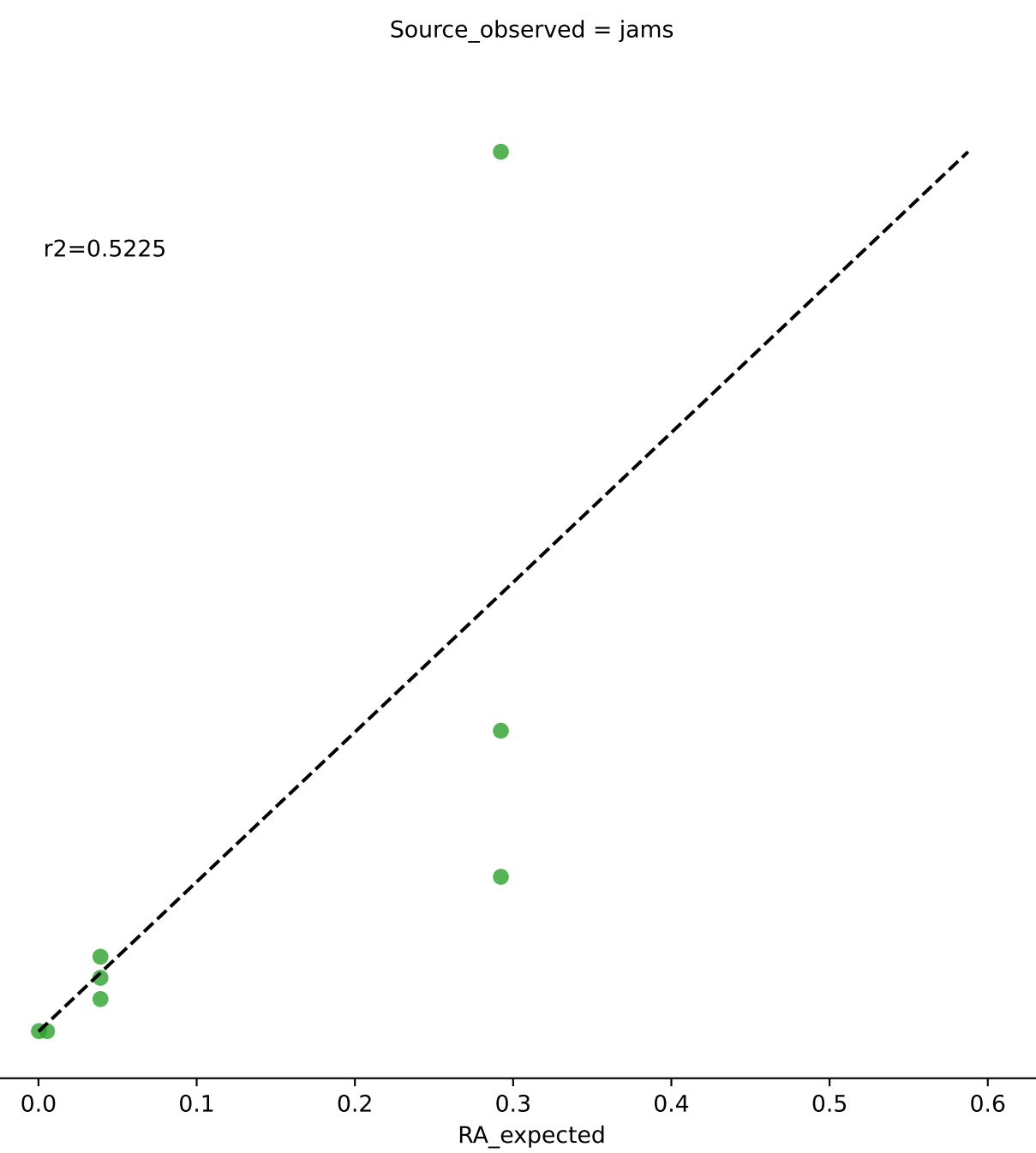
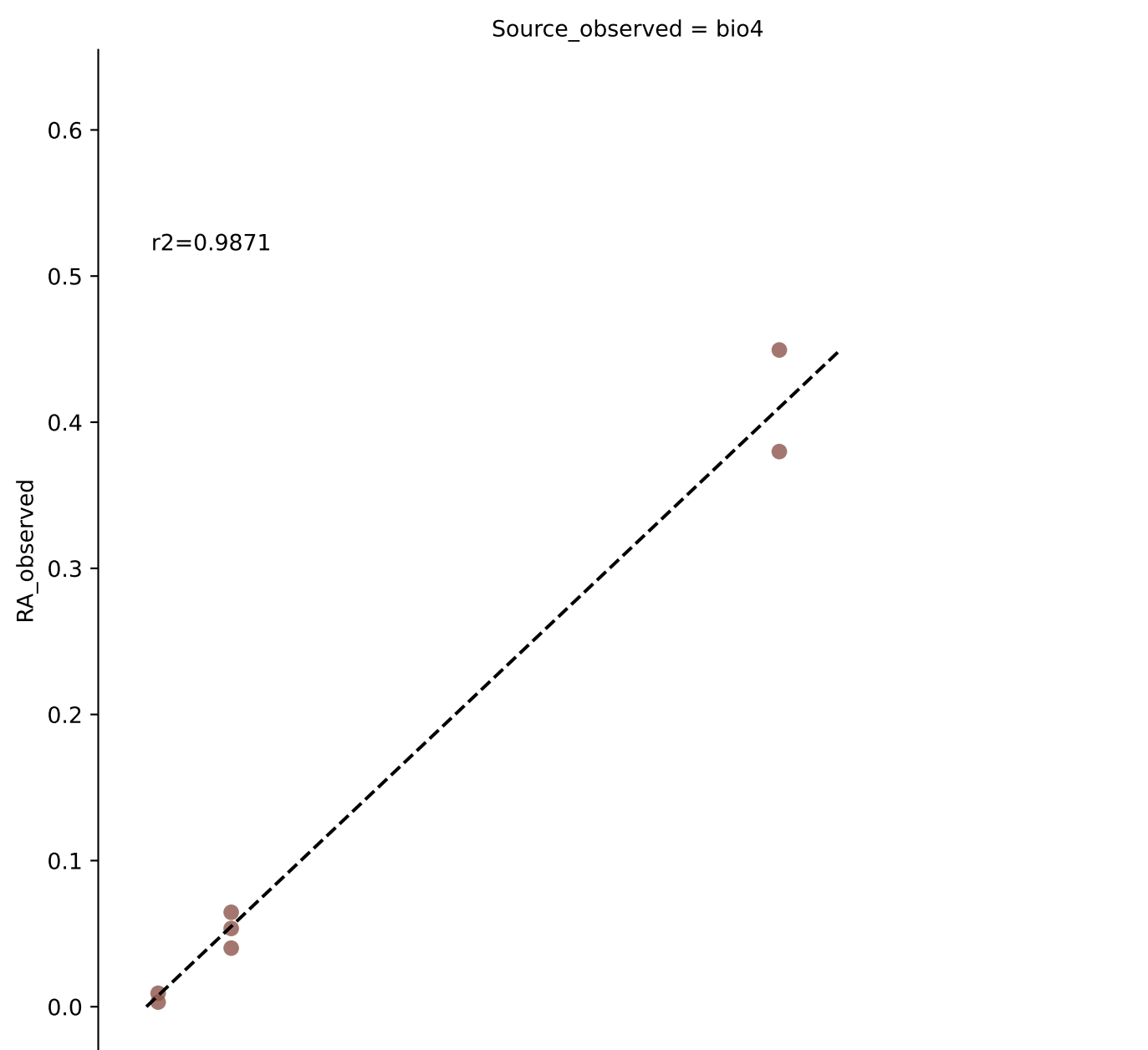


Bivariate Linear Regression for Sample MIX-B in Experiment nist (Genus at filter threshold 0.0001)

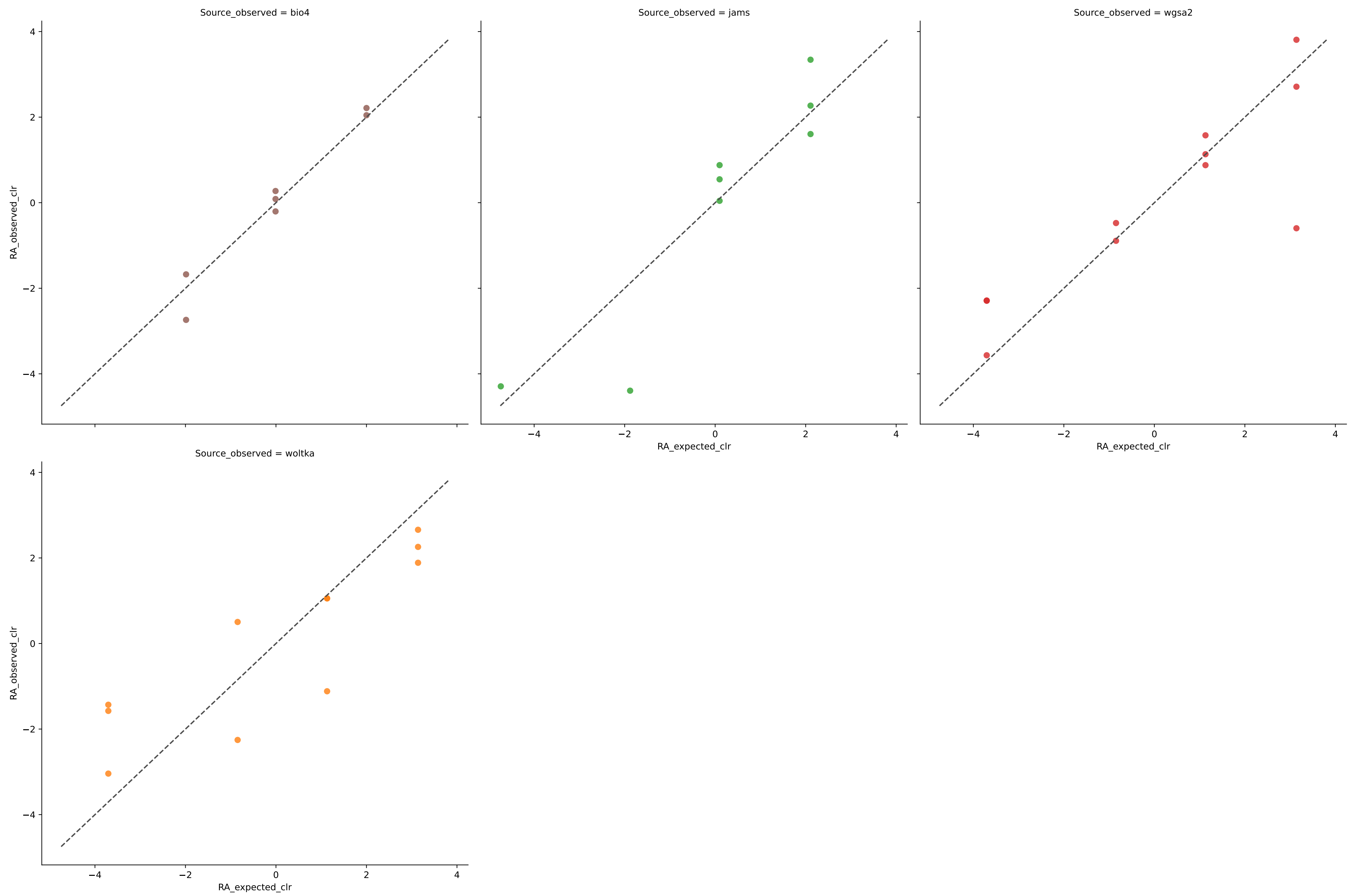


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	8	0.6977	0.0532	4.7646	0.7871	0.0857	100.0000	0.0000
jams	8	0.9577	0.0271	4.7510	0.8917	0.0345	100.0000	0.0000
wgsa2	11	0.9691	0.0167	5.2327	0.9083	0.0241	100.0000	0.0000
woltka	11	0.7965	0.0435	6.1353	0.7607	0.0631	100.0000	0.0000

Bivariate Linear Regression for Sample MIX-C in Experiment nist (Genus at filter threshold 0.0001)

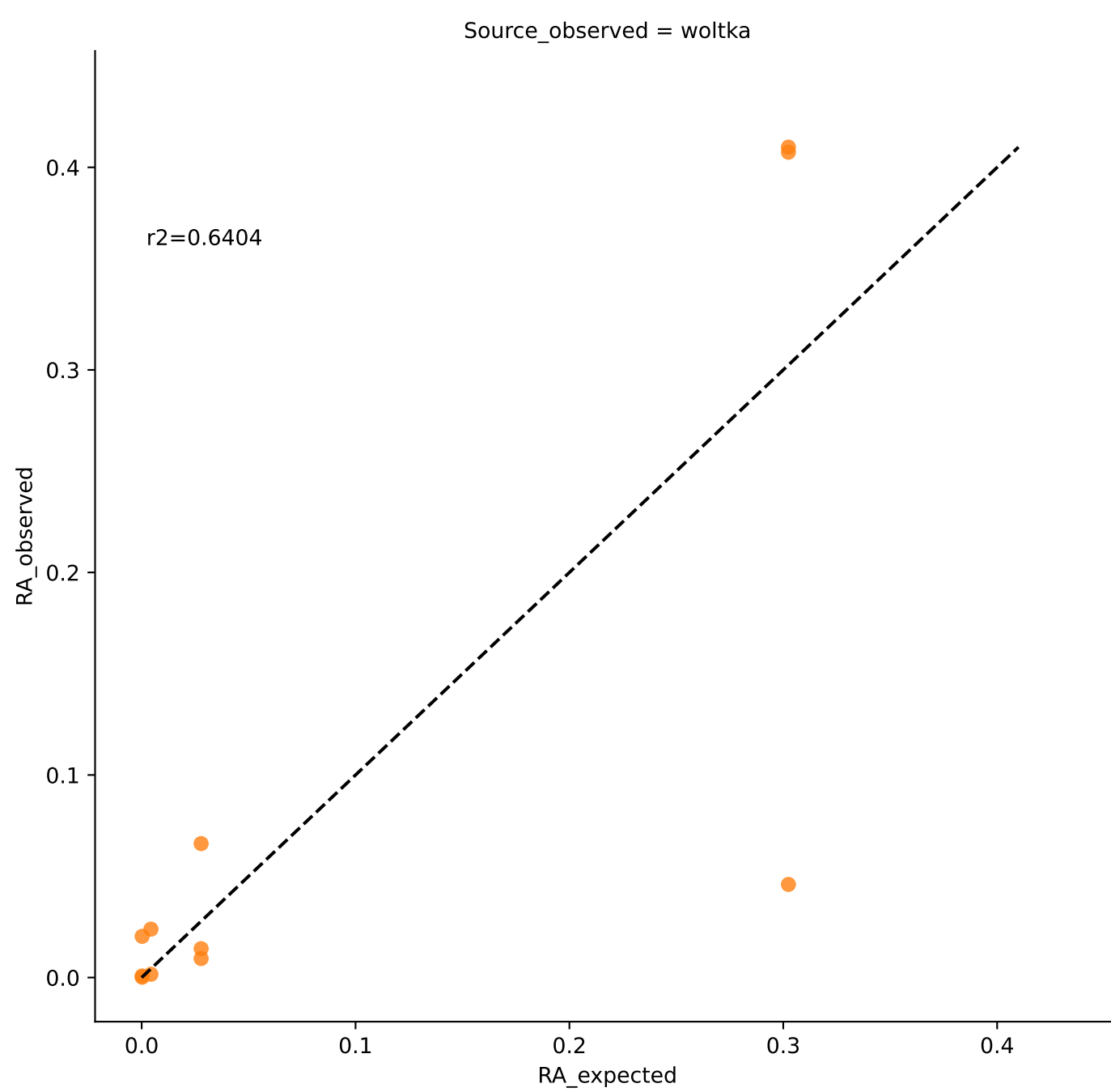
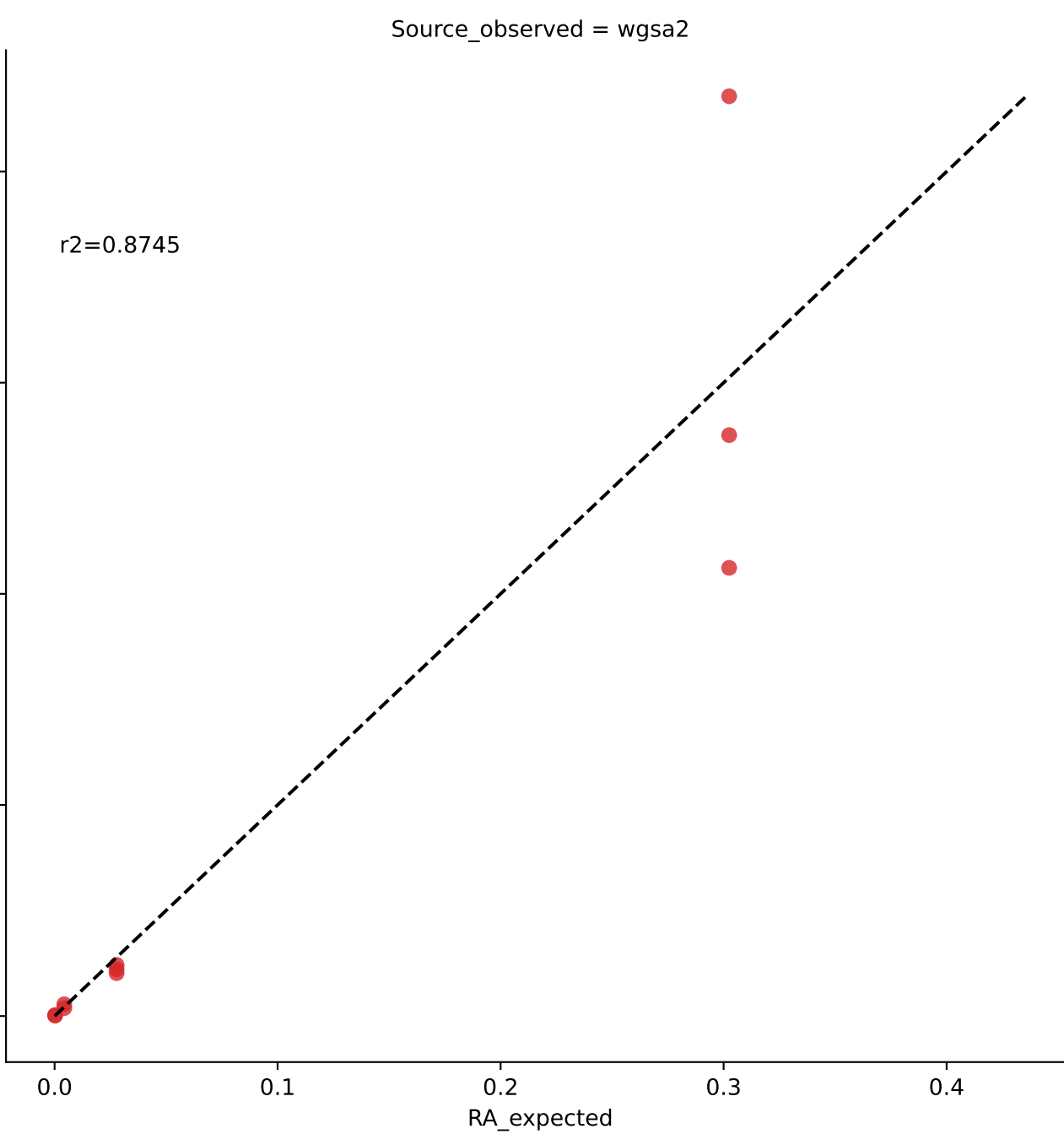
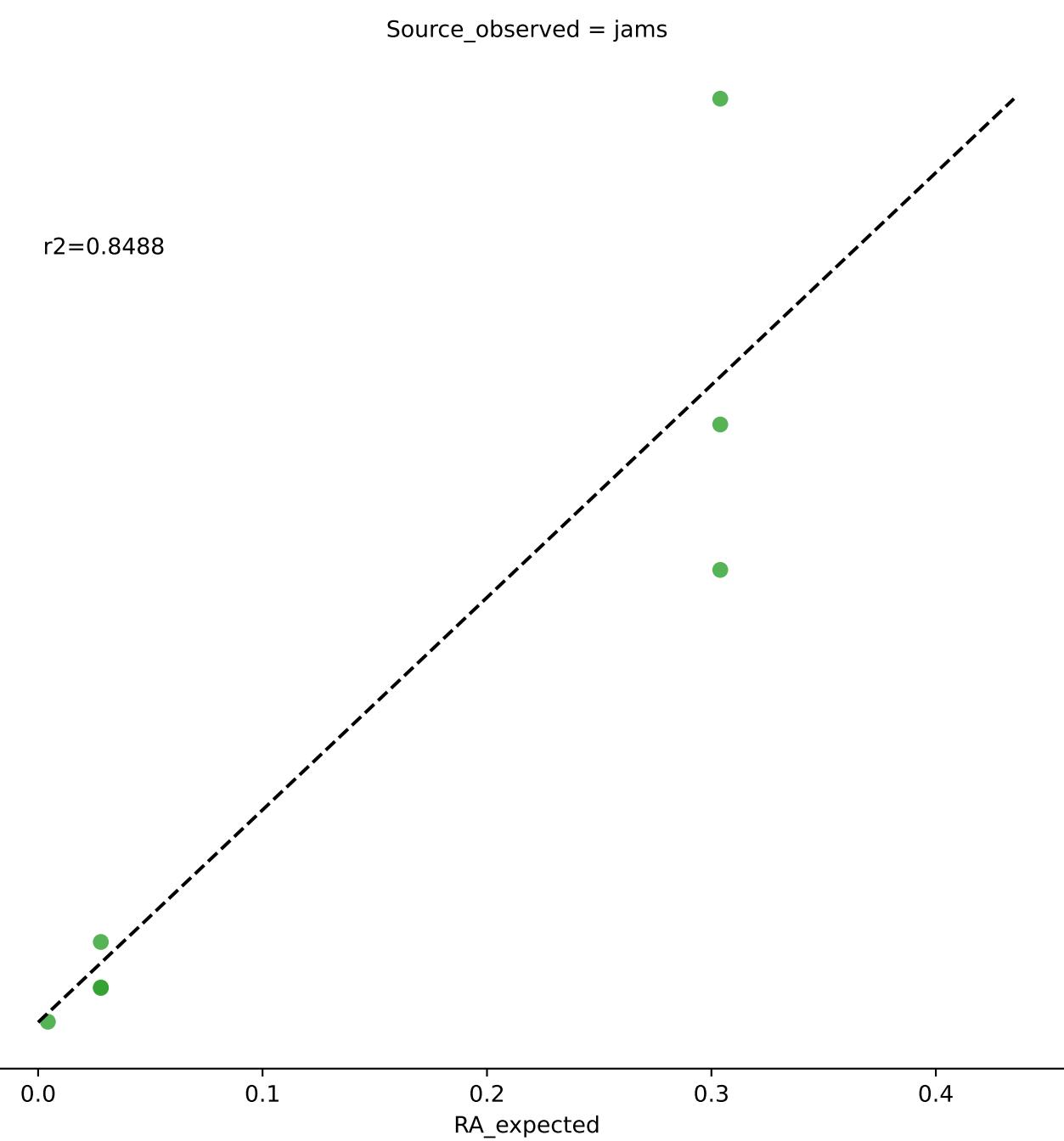
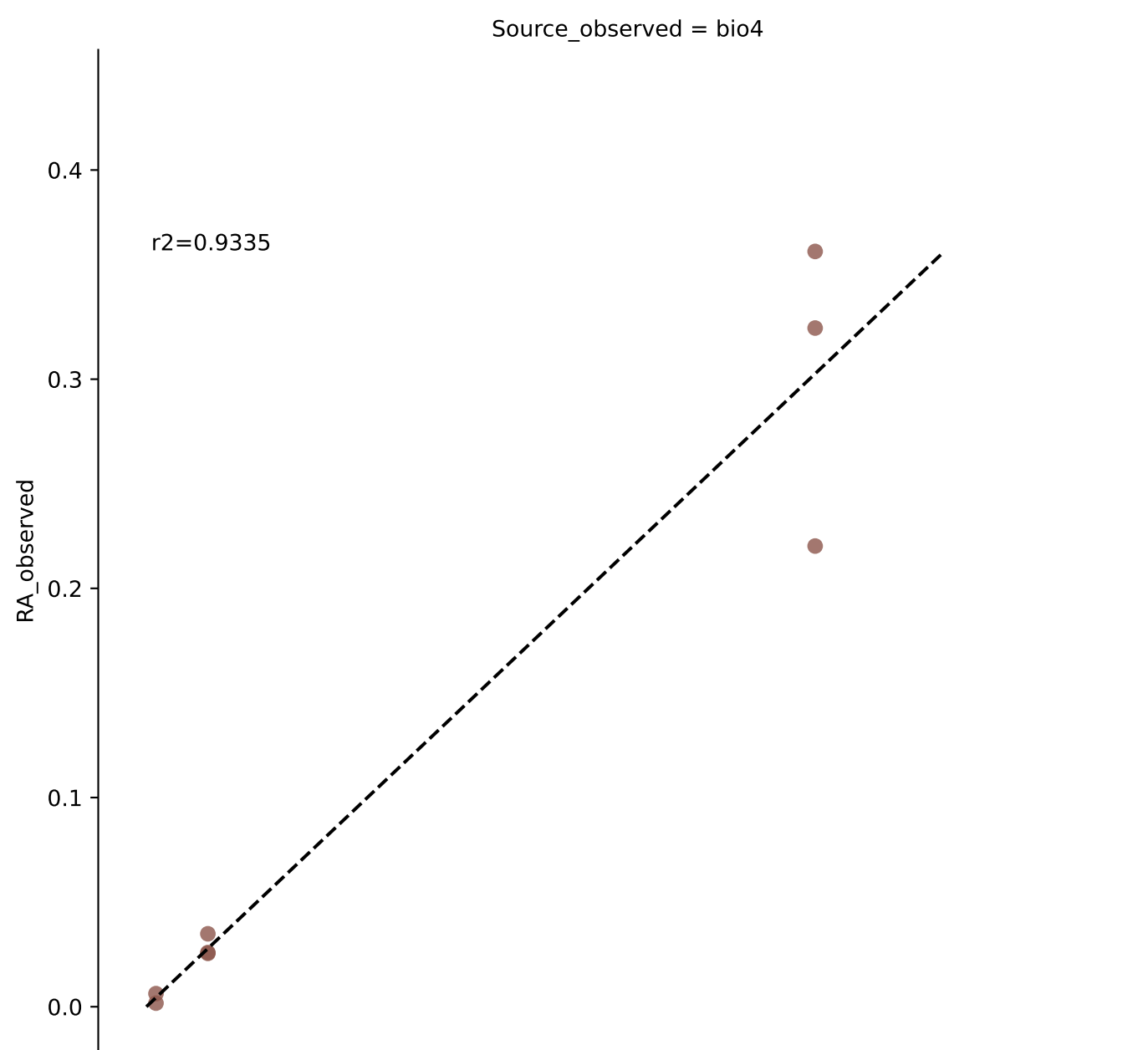


Bivariate Linear Regression for Sample MIX-C in Experiment nist (Genus at filter threshold 0.0001)

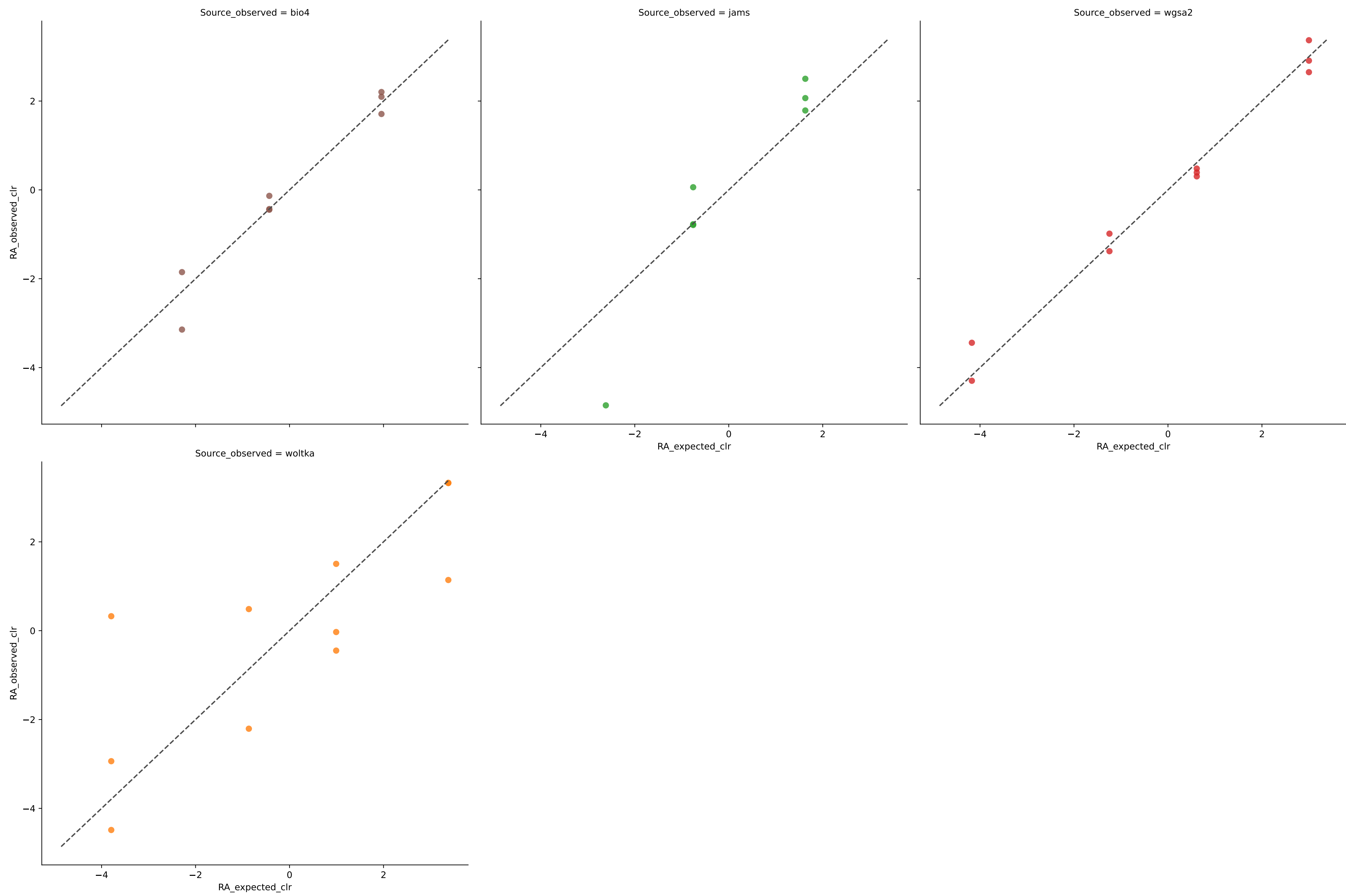


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	7	0.9871	0.0145	0.9149	0.9493	0.0200	100.0000	0.0000
jams	8	0.5225	0.0765	3.0229	0.6939	0.1283	100.0000	0.0000
wgsa2	11	0.4320	0.0674	4.3618	0.6292	0.1344	100.0000	0.0000
woltka	11	0.8372	0.0358	4.6458	0.8029	0.0498	100.0000	0.0000

Bivariate Linear Regression for Sample MIX-D in Experiment nist (Genus at filter threshold 0.0001)

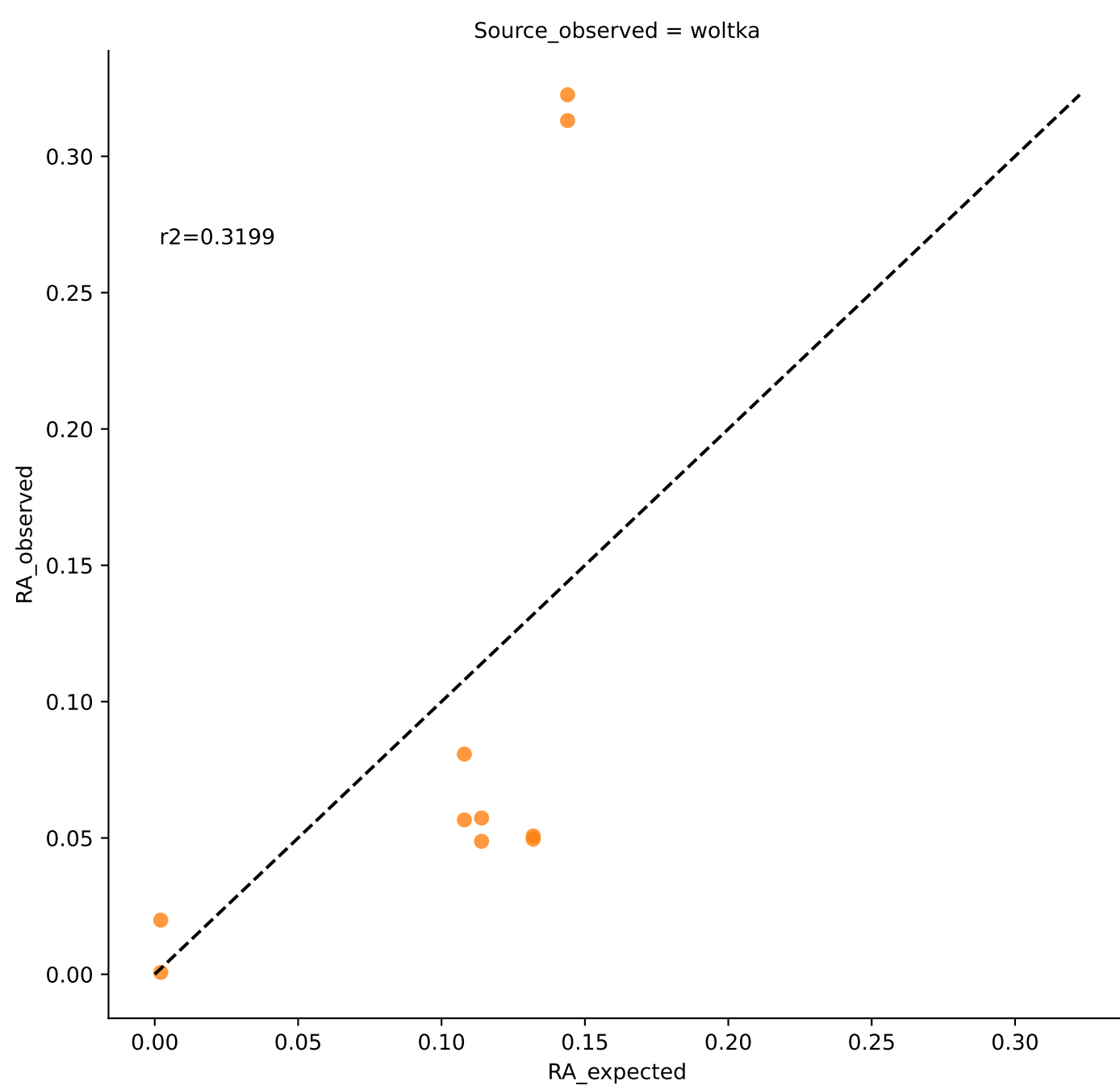
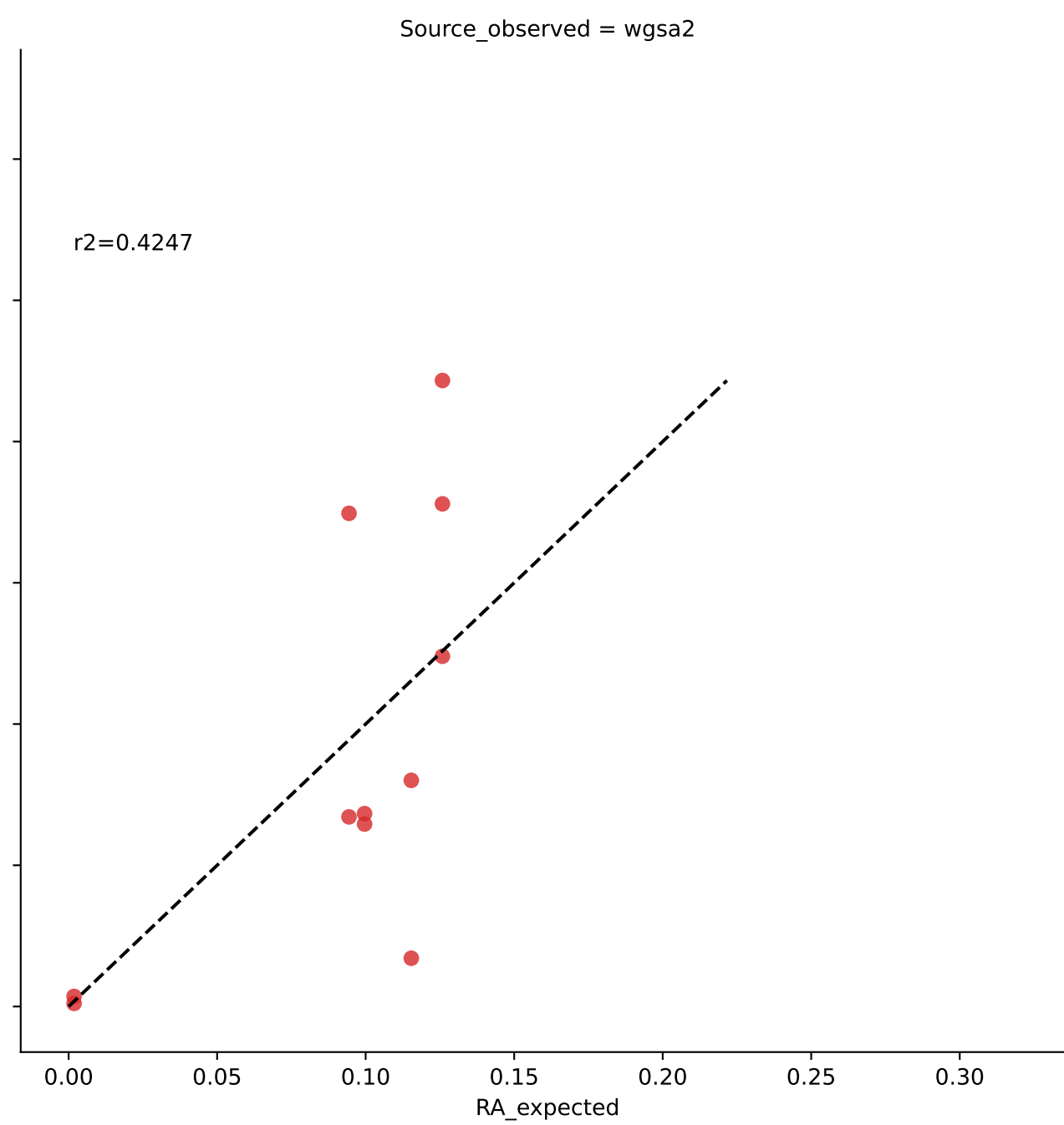
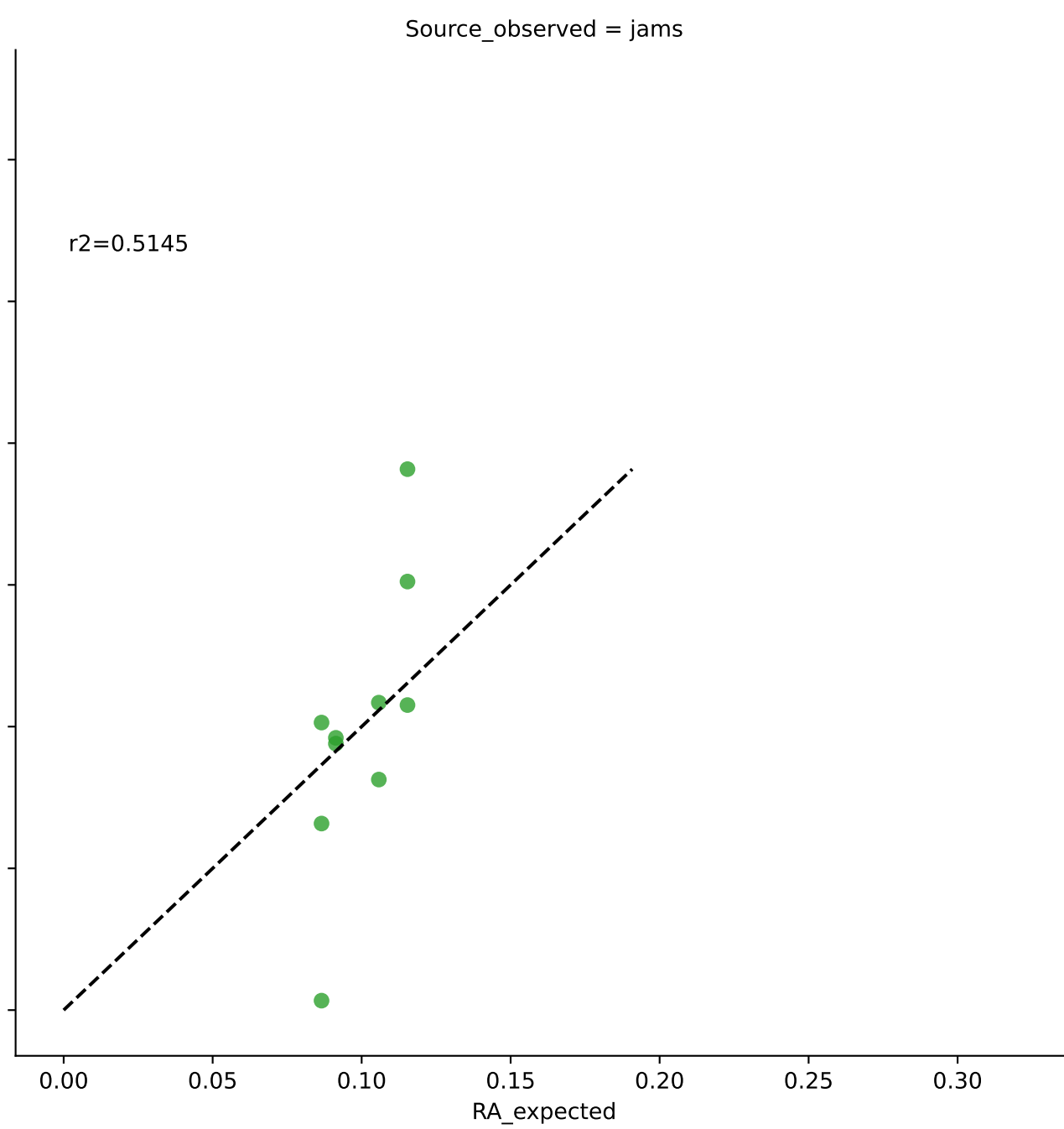
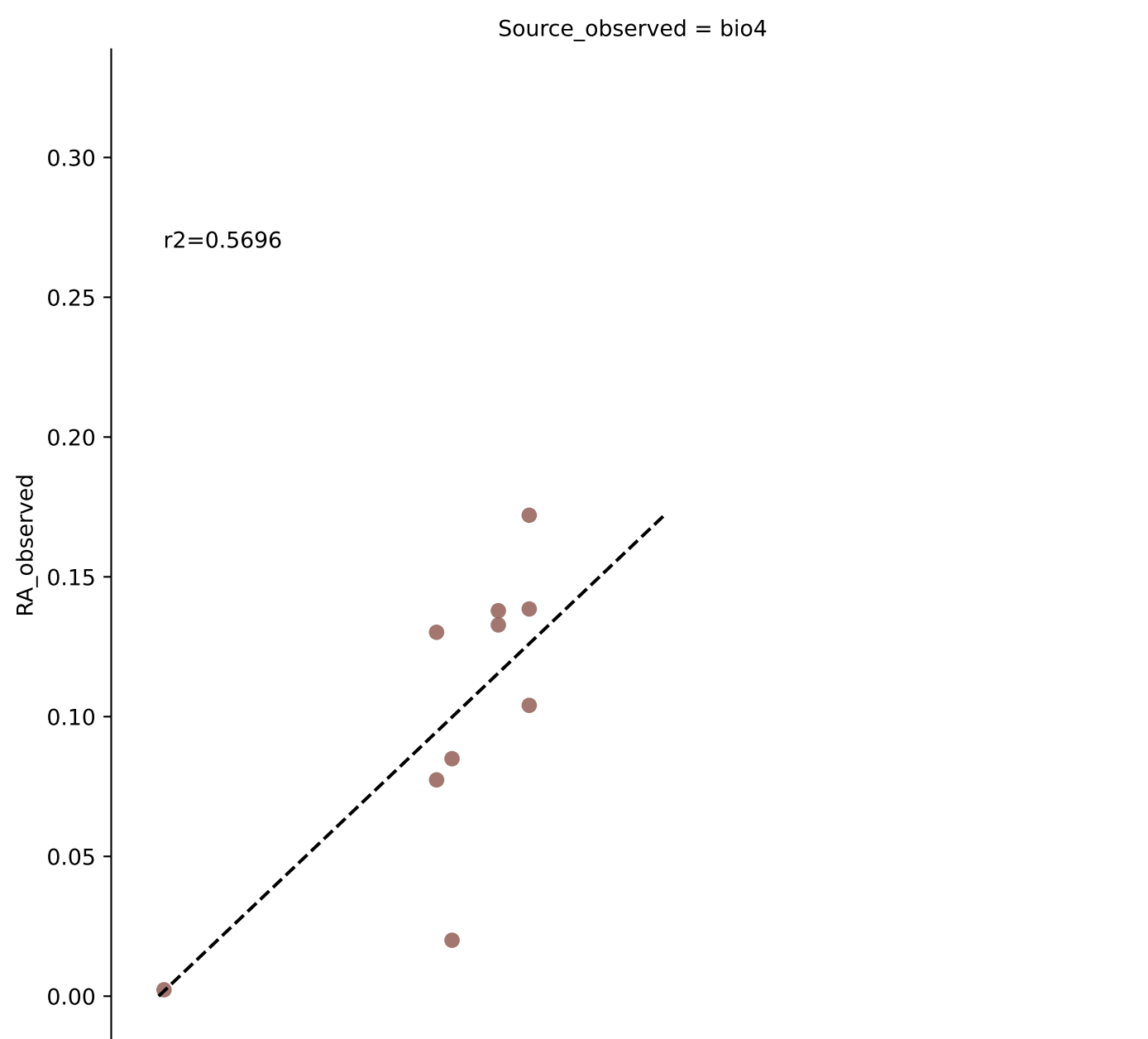


Bivariate Linear Regression for Sample MIX-D in Experiment nist (Genus at filter threshold 0.0001)

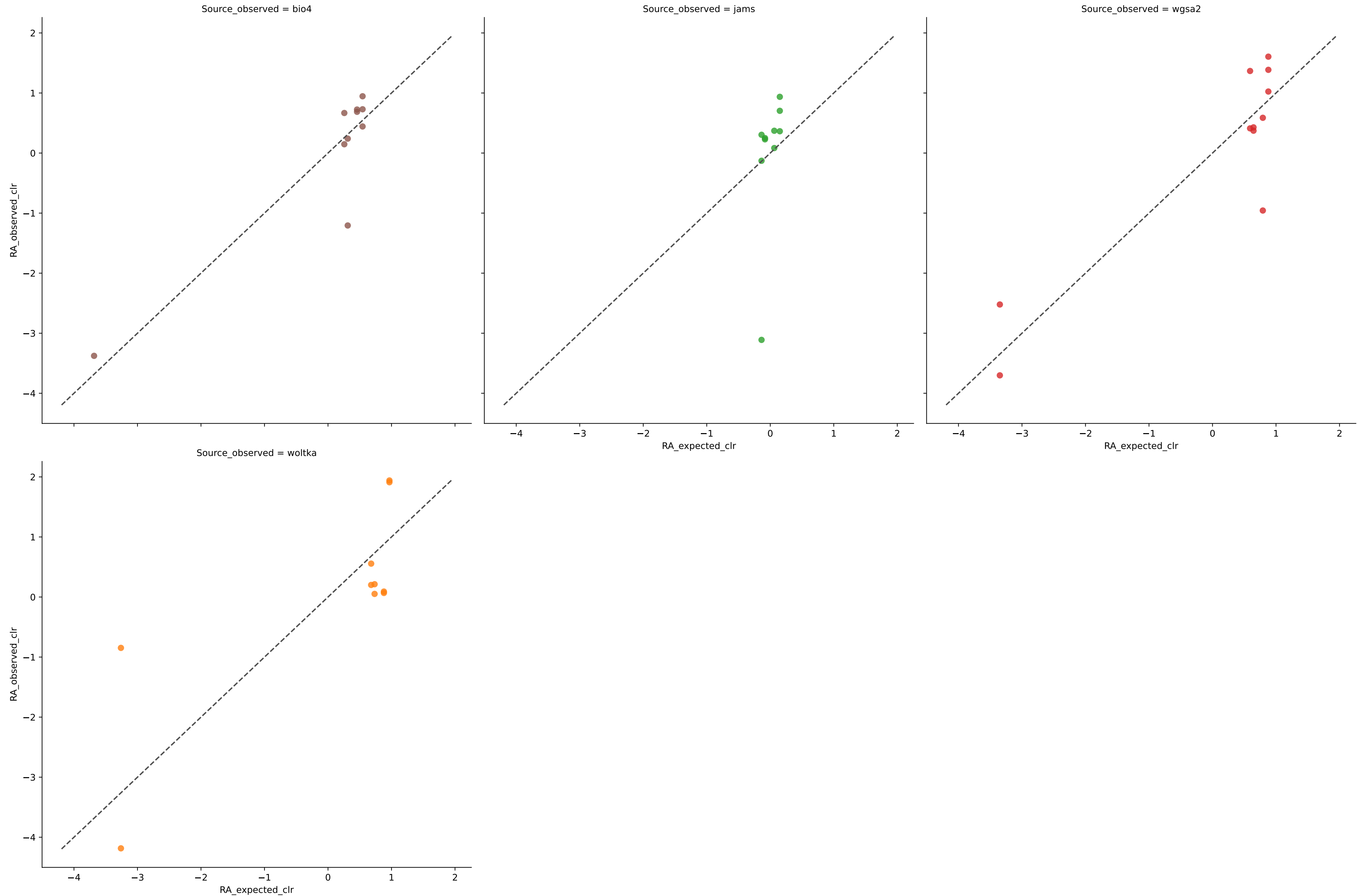


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	8	0.9335	0.0223	1.0739	0.9107	0.0367	100.0000	0.0000
jams	7	0.8488	0.0402	2.5764	0.8592	0.0613	100.0000	0.0000
wgsa2	10	0.8745	0.0269	1.0323	0.8653	0.0517	100.0000	0.0000
woltka	11	0.6404	0.0529	5.4967	0.7088	0.0910	100.0000	0.0000

Bivariate Linear Regression for Sample EG in Experiment nist (Species at filter threshold 0.0001)

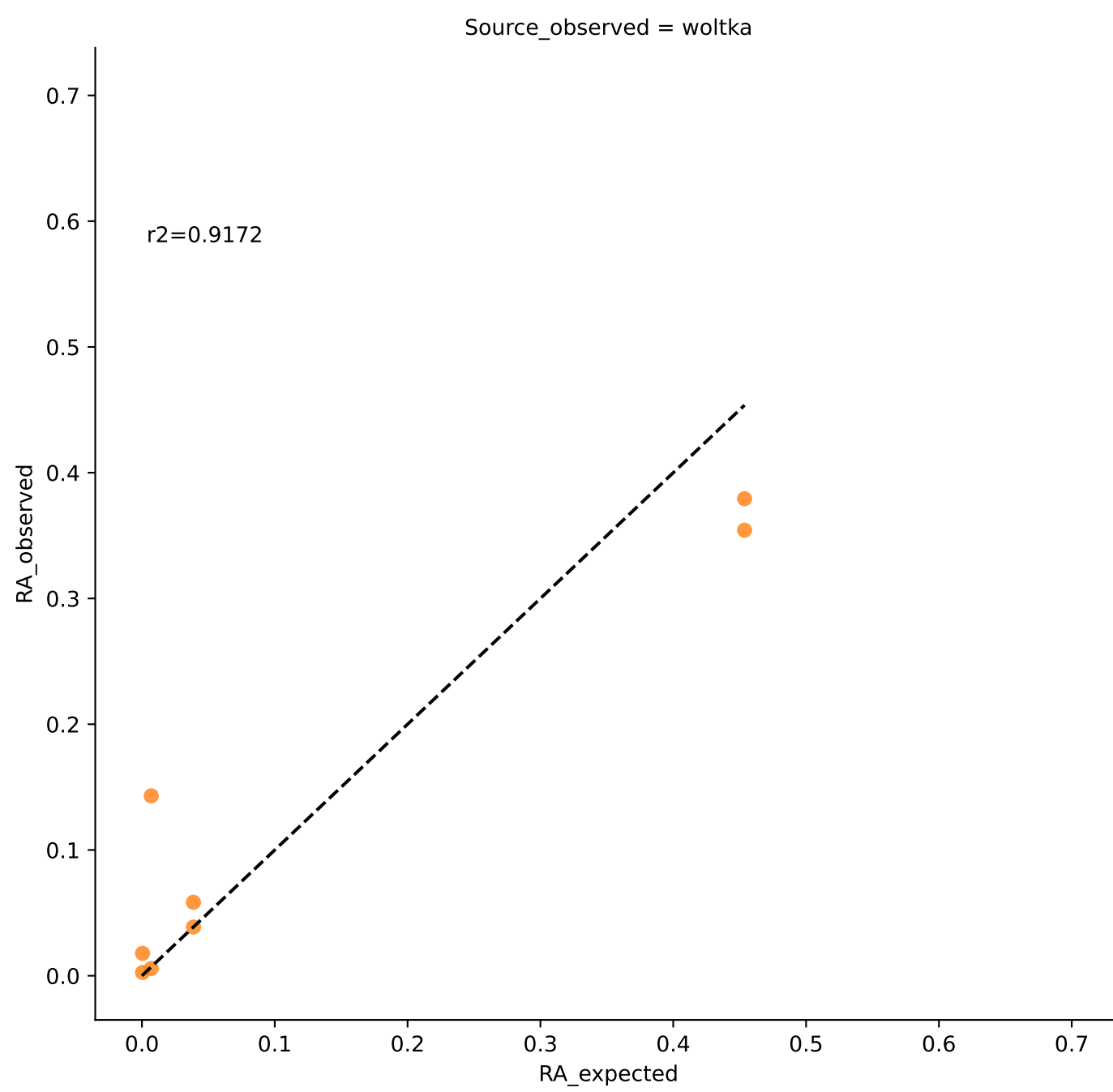
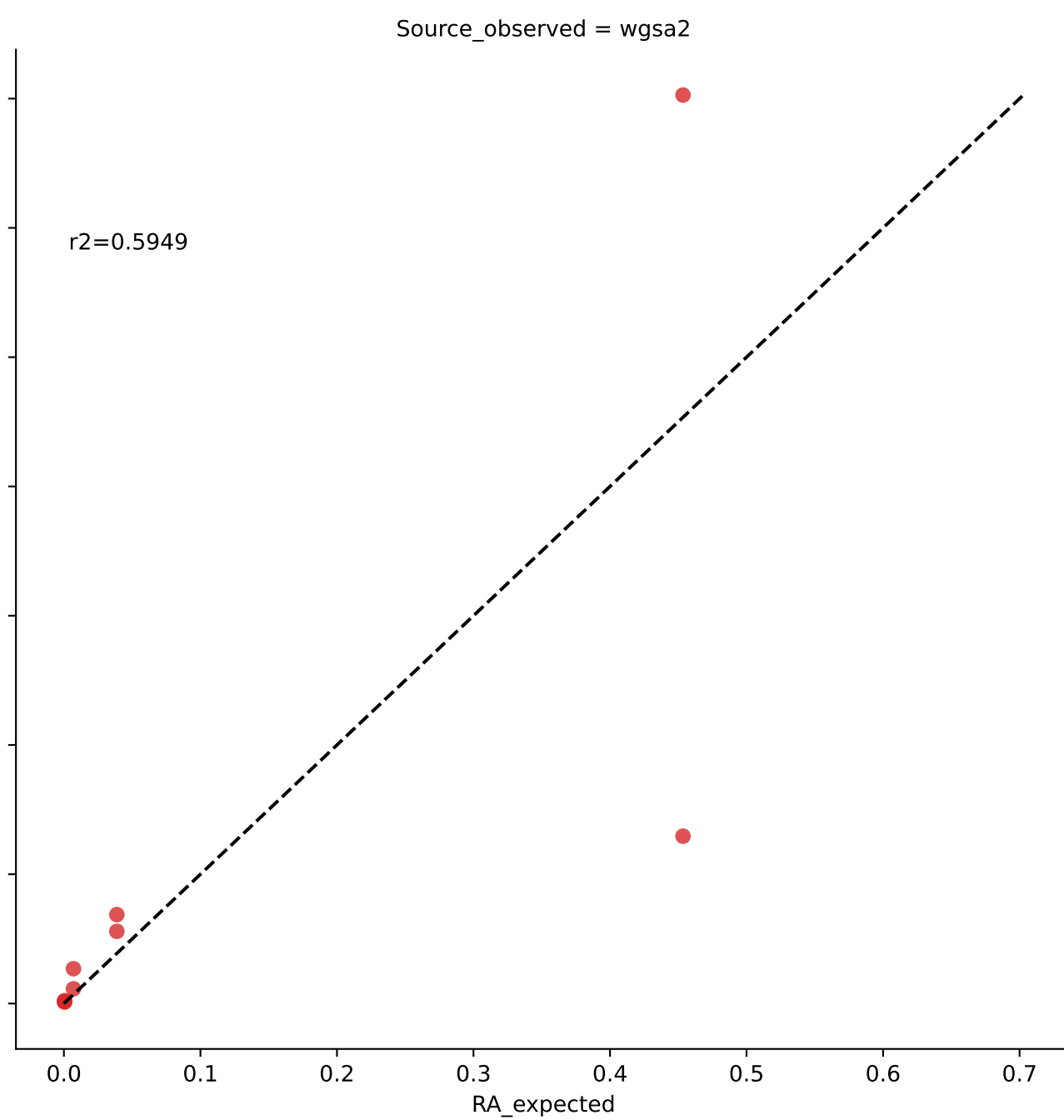
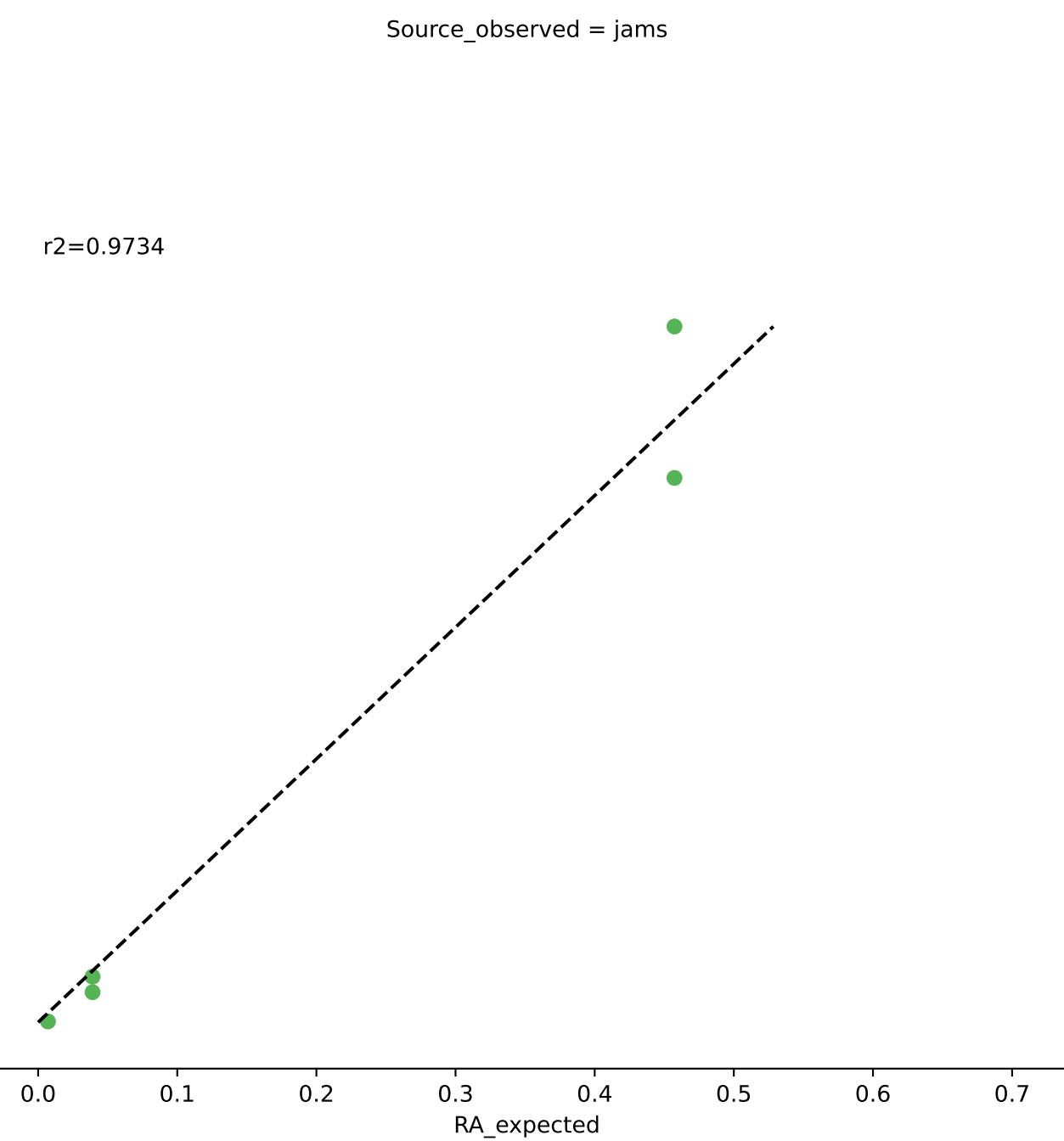
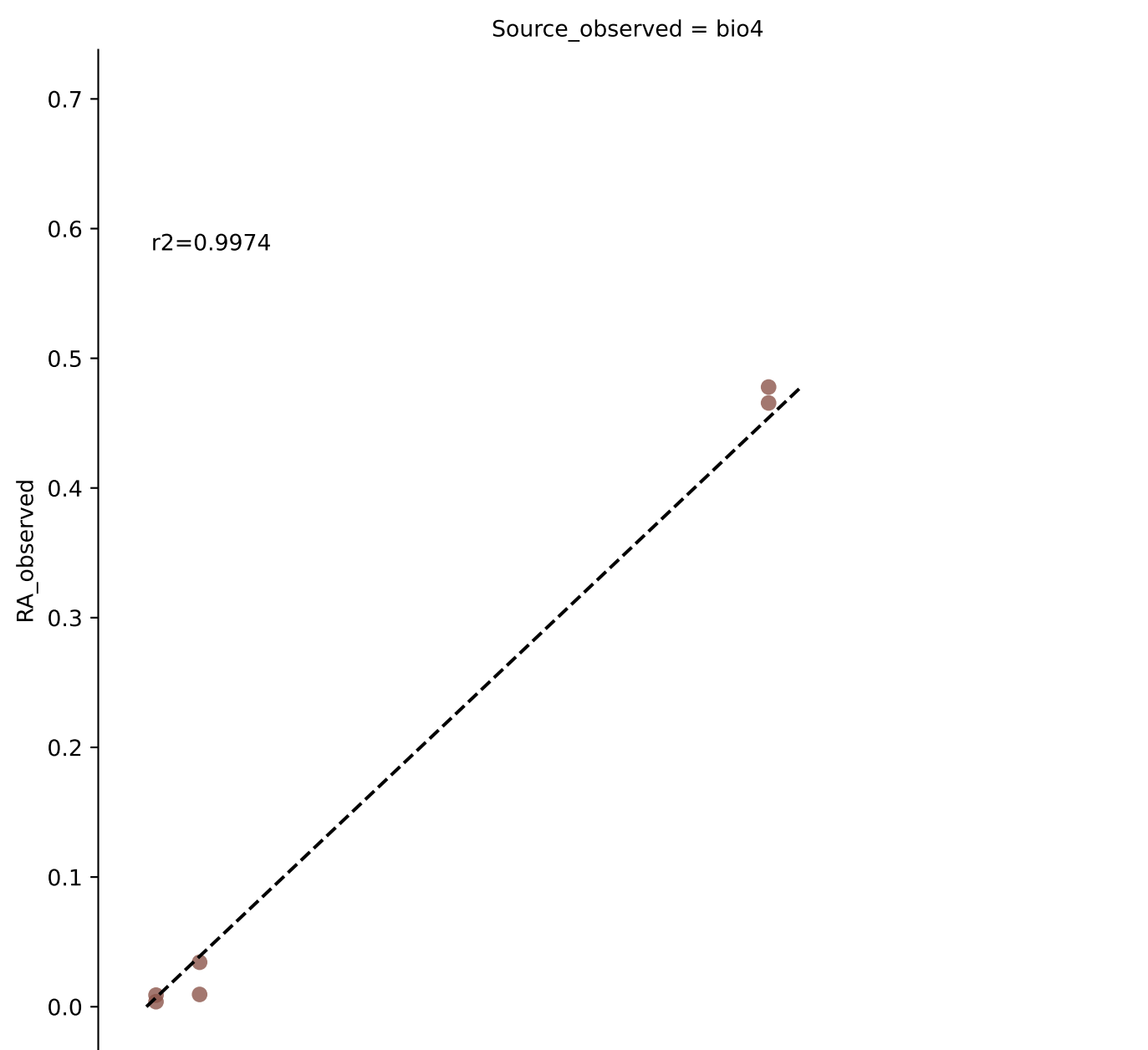


Bivariate Linear Regression for Sample EG in Experiment nist (Species at filter threshold 0.0001)

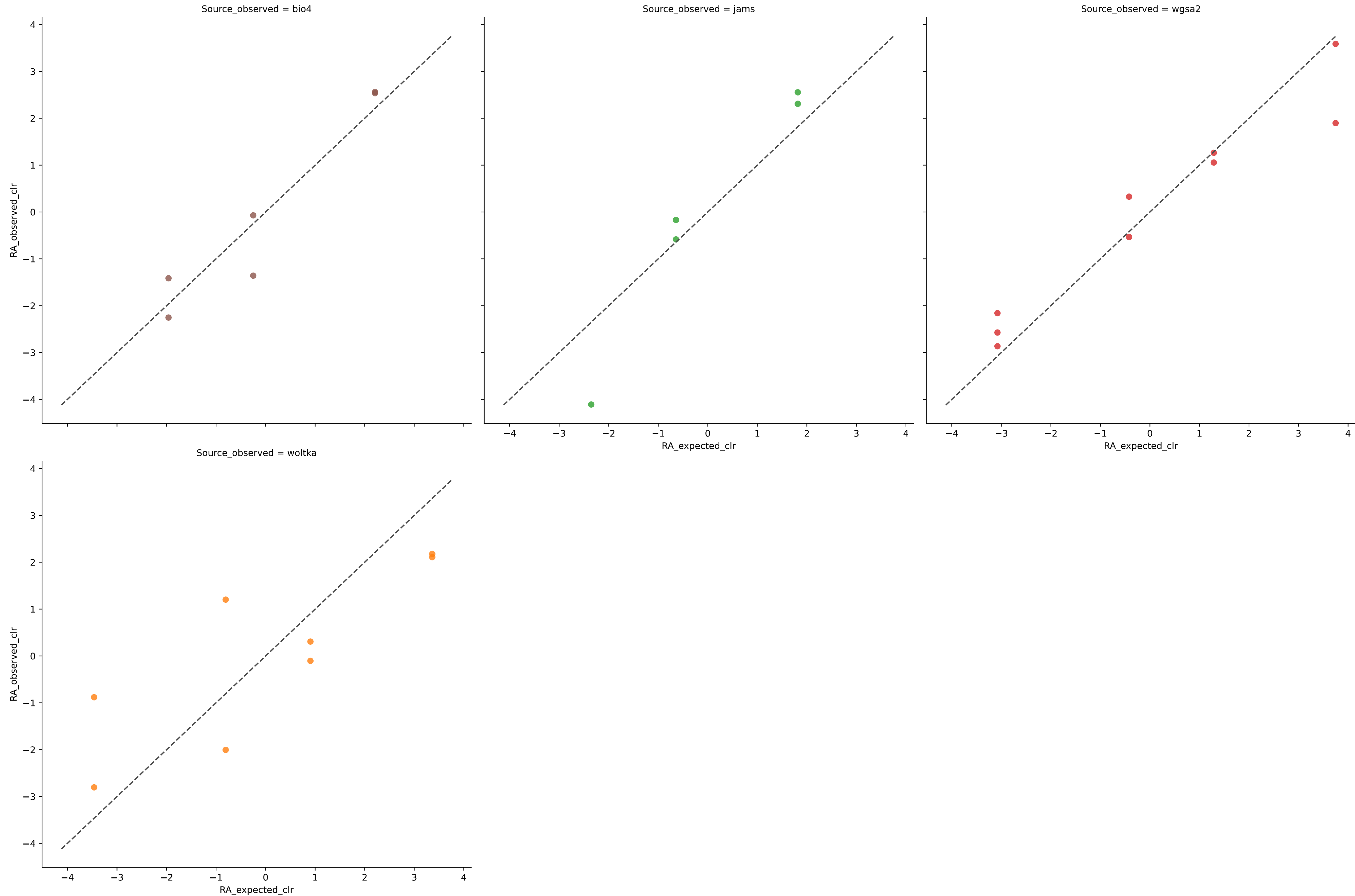


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	10	0.5696	0.0268	1.7058	0.8661	0.0342	100.0000	0.0000
jams	10	0.5145	0.0272	3.2110	0.8638	0.0390	100.0000	0.0000
wgsa2	11	0.4247	0.0418	2.3389	0.7702	0.0541	100.0000	0.0000
woltka	10	0.3199	0.0731	3.2836	0.6345	0.0922	100.0000	0.0000

Bivariate Linear Regression for Sample MIX-A in Experiment nist (Species at filter threshold 0.0001)

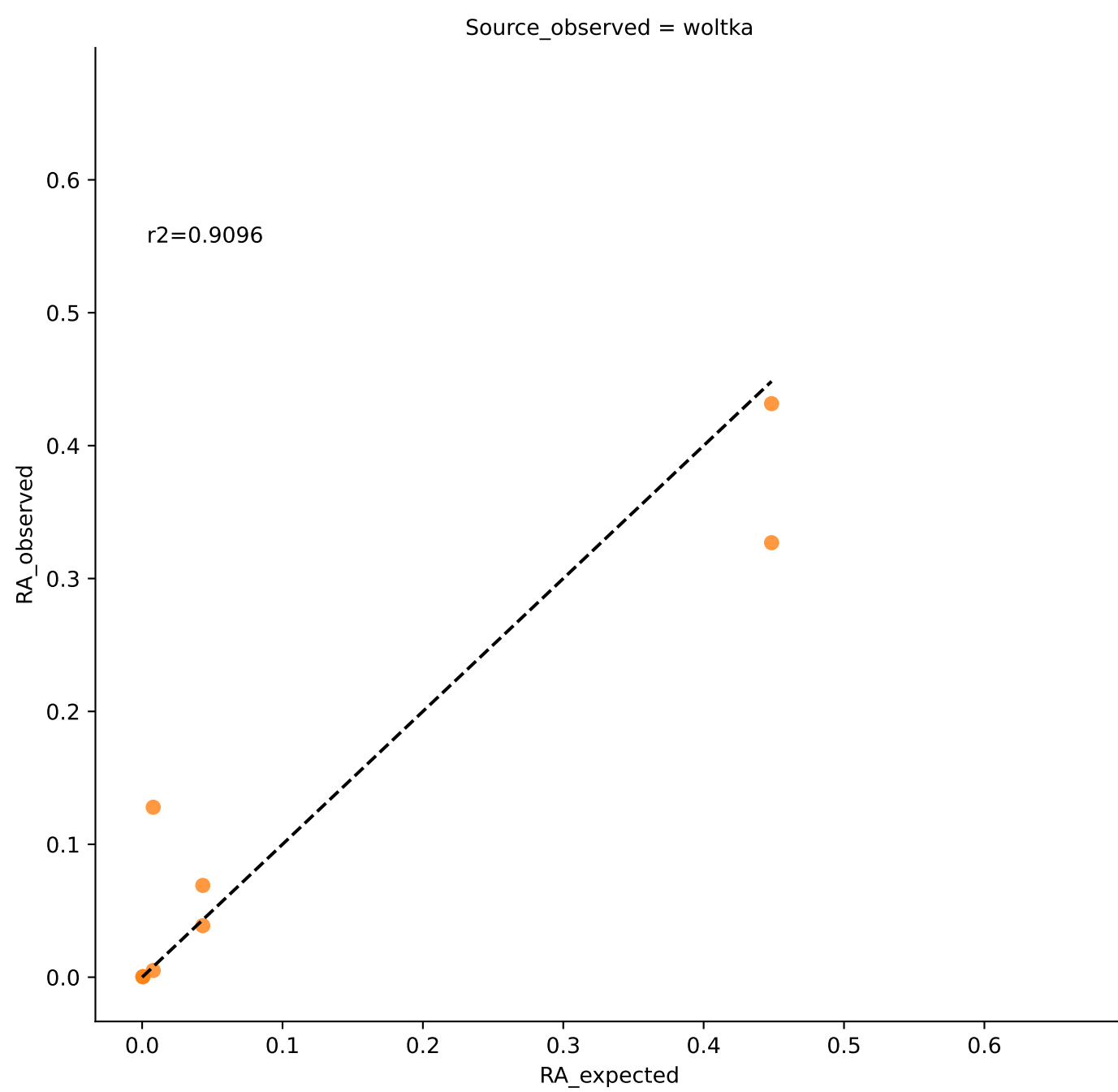
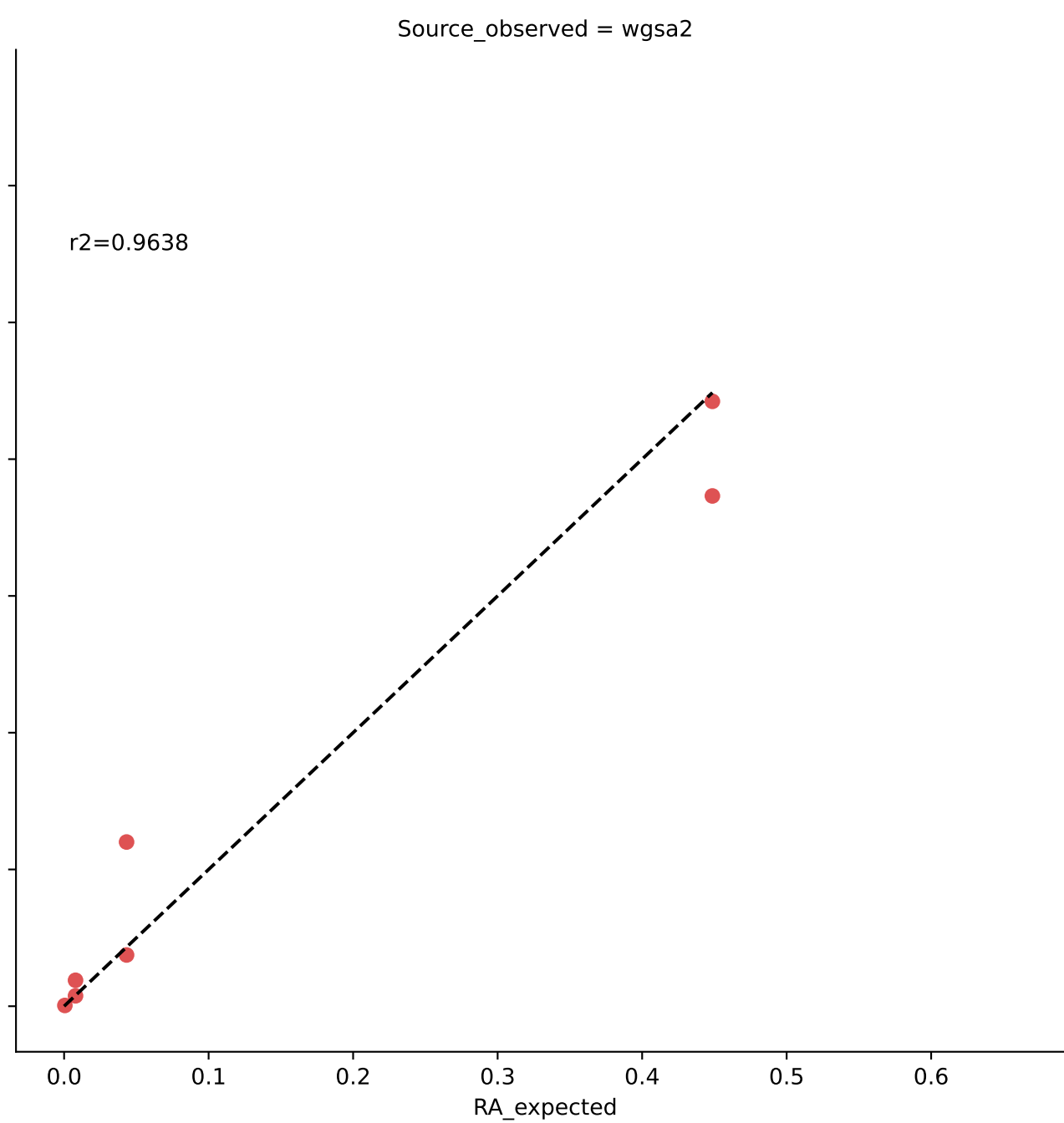
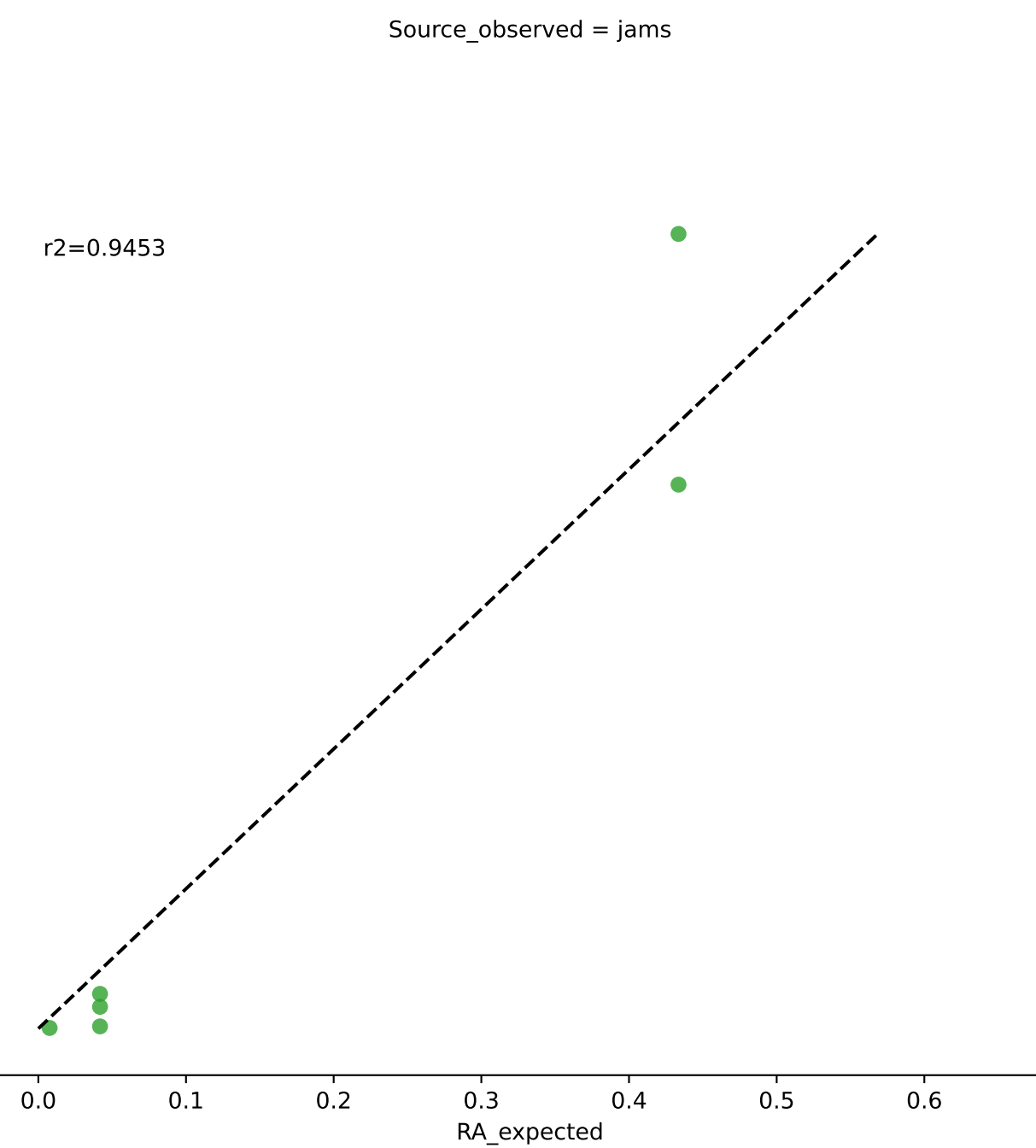
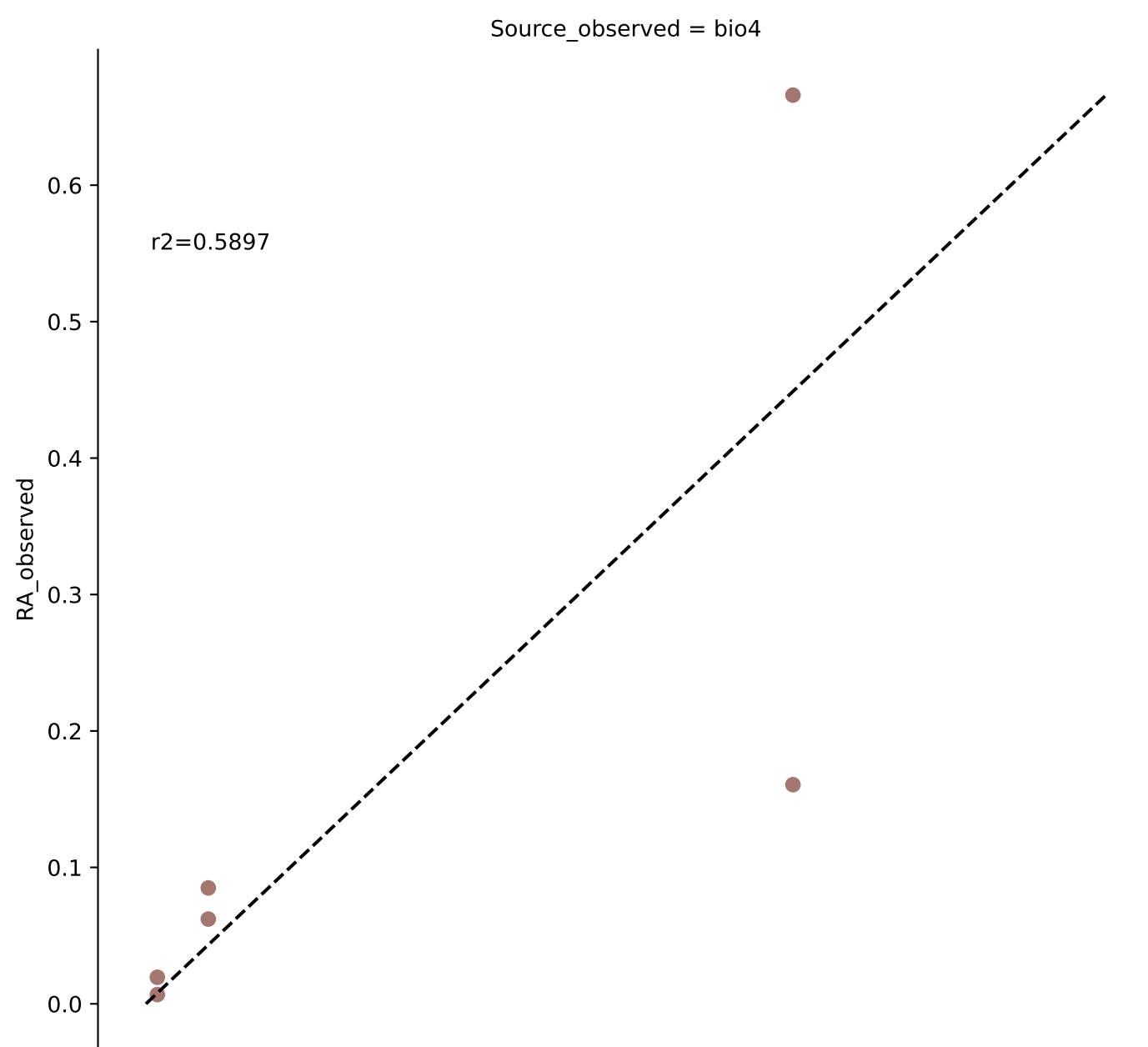


Bivariate Linear Regression for Sample MIX-A in Experiment nist (Species at filter threshold 0.0001)

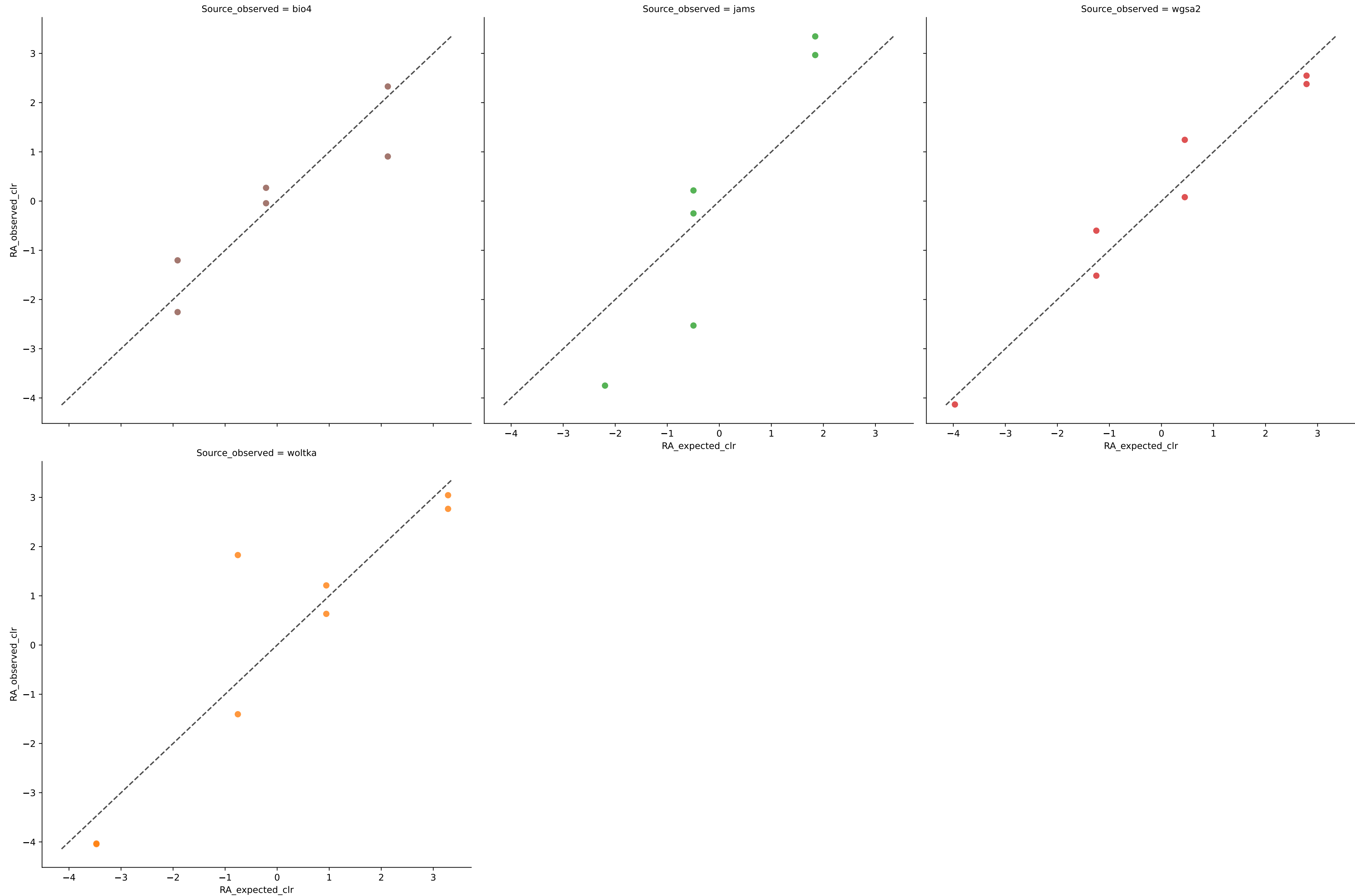


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	6	0.9974	0.0124	1.3695	0.9629	0.0163	100.0000	0.0000
jams	5	0.9734	0.0284	2.0241	0.9290	0.0382	100.0000	0.0000
wgsa2	9	0.5949	0.0720	2.2860	0.6760	0.1369	100.0000	0.0000
woltka	8	0.9172	0.0438	4.1130	0.8249	0.0658	100.0000	0.0000

Bivariate Linear Regression for Sample MIX-B in Experiment nist (Species at filter threshold 0.0001)

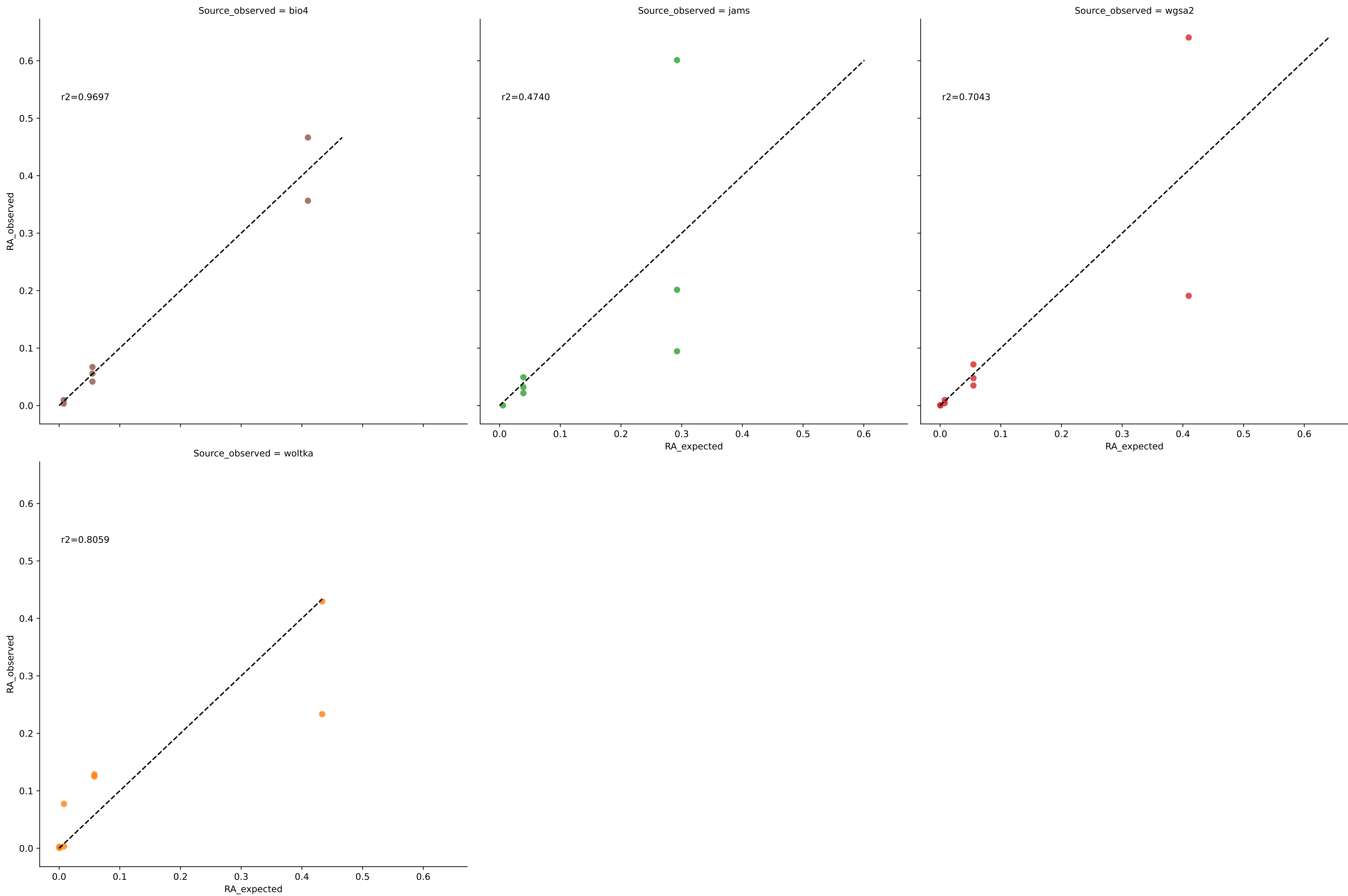


Bivariate Linear Regression for Sample MIX-B in Experiment nist (Species at filter threshold 0.0001)

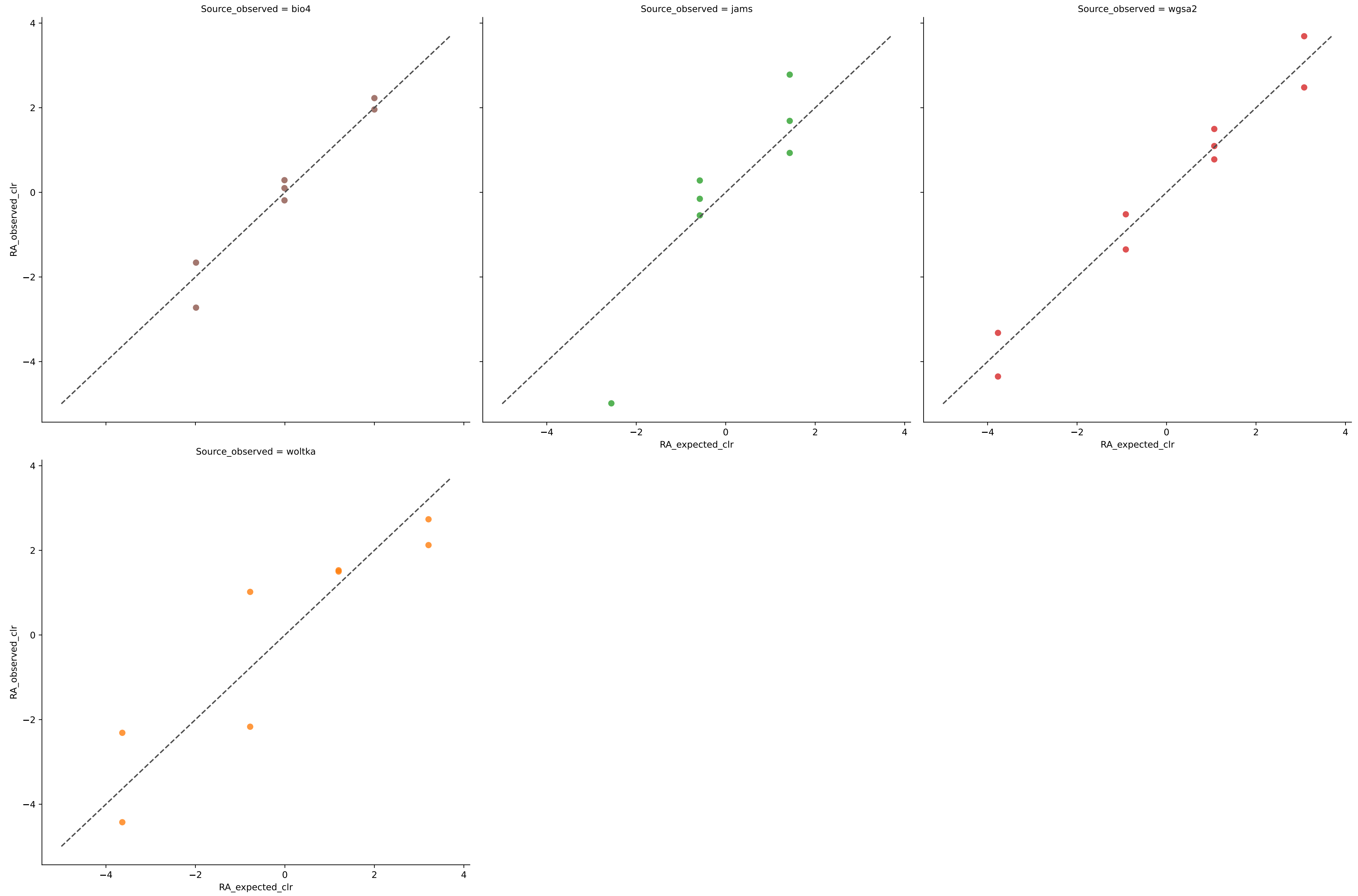


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	6	0.5897	0.0964	1.5530	0.7107	0.1486	100.0000	0.0000
jams	6	0.9453	0.0450	3.2599	0.8651	0.0616	100.0000	0.0000
wgsa2	7	0.9638	0.0251	1.2308	0.9121	0.0411	100.0000	0.0000
woltka	8	0.9096	0.0365	2.8687	0.8542	0.0614	100.0000	0.0000

Bivariate Linear Regression for Sample MIX-C in Experiment nist (Species at filter threshold 0.0001)

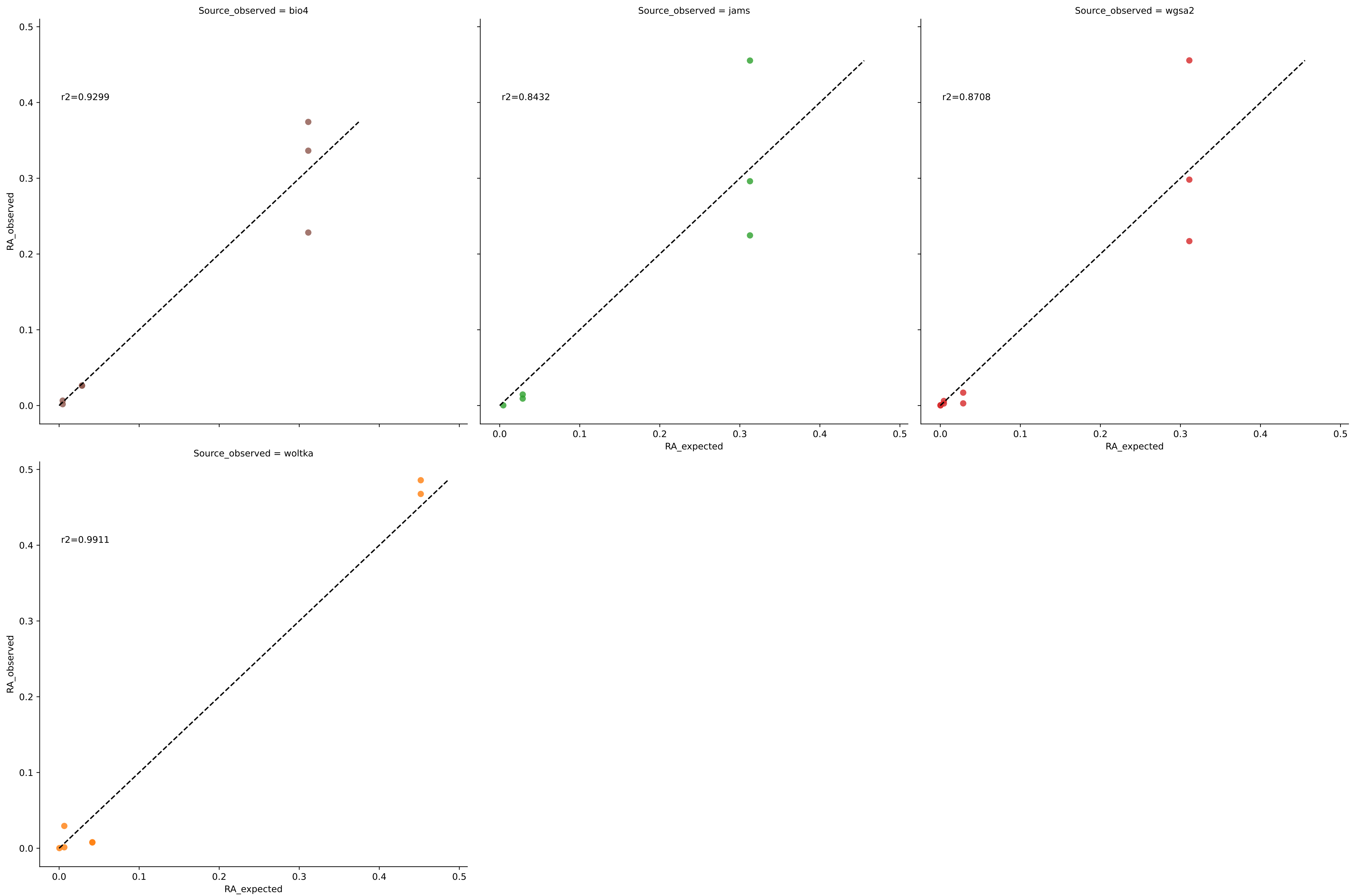


Bivariate Linear Regression for Sample MIX-C in Experiment nist (Species at filter threshold 0.0001)

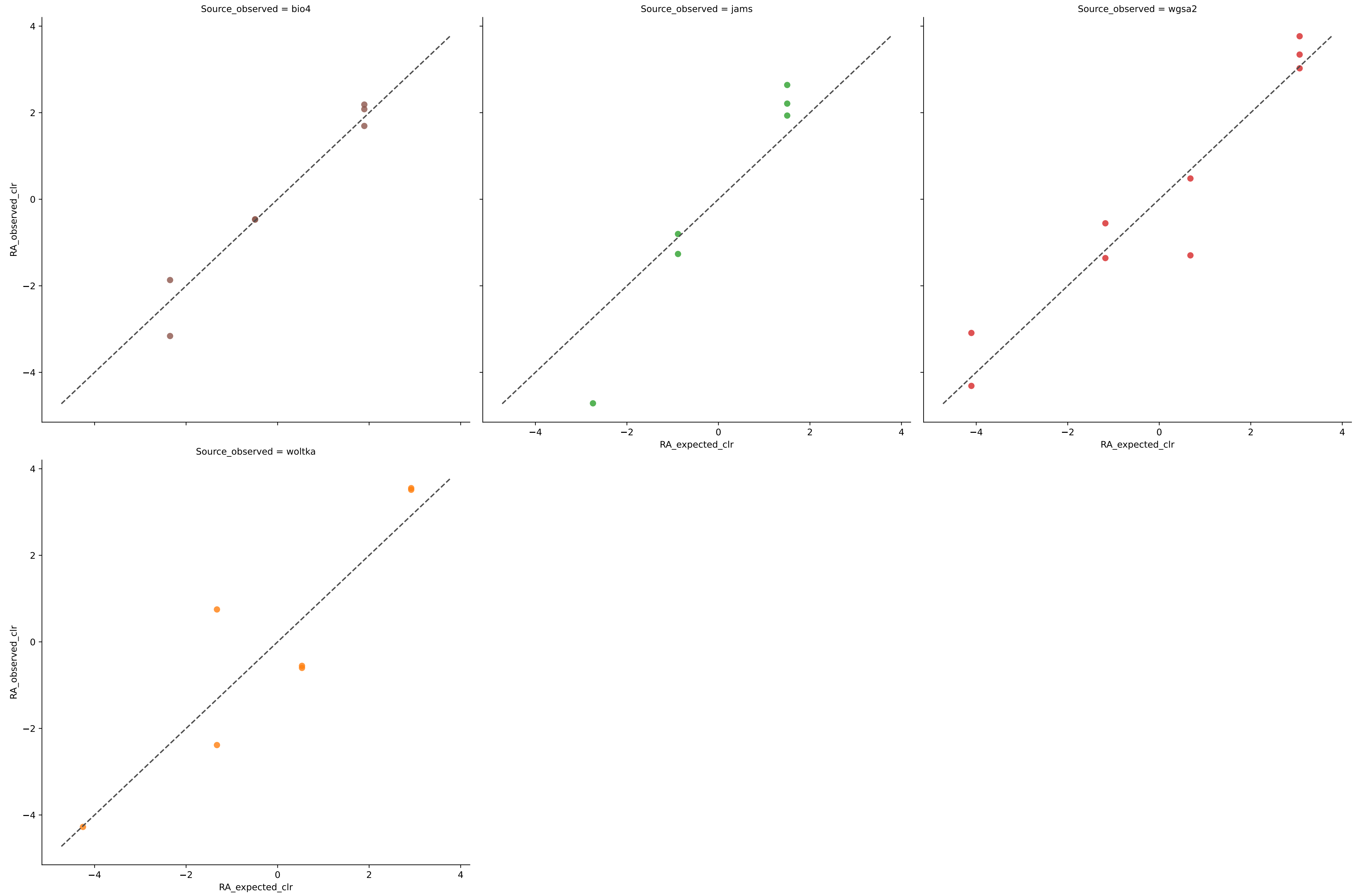


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	7	0.9697	0.0203	0.9148	0.9288	0.0302	100.0000	0.0000
jams	7	0.4740	0.0911	2.9945	0.6812	0.1430	100.0000	0.0000
wgsa2	9	0.7043	0.0555	1.3745	0.7505	0.1064	100.0000	0.0000
woltka	8	0.8059	0.0522	3.0224	0.7911	0.0824	100.0000	0.0000

Bivariate Linear Regression for Sample MIX-D in Experiment nist (Species at filter threshold 0.0001)

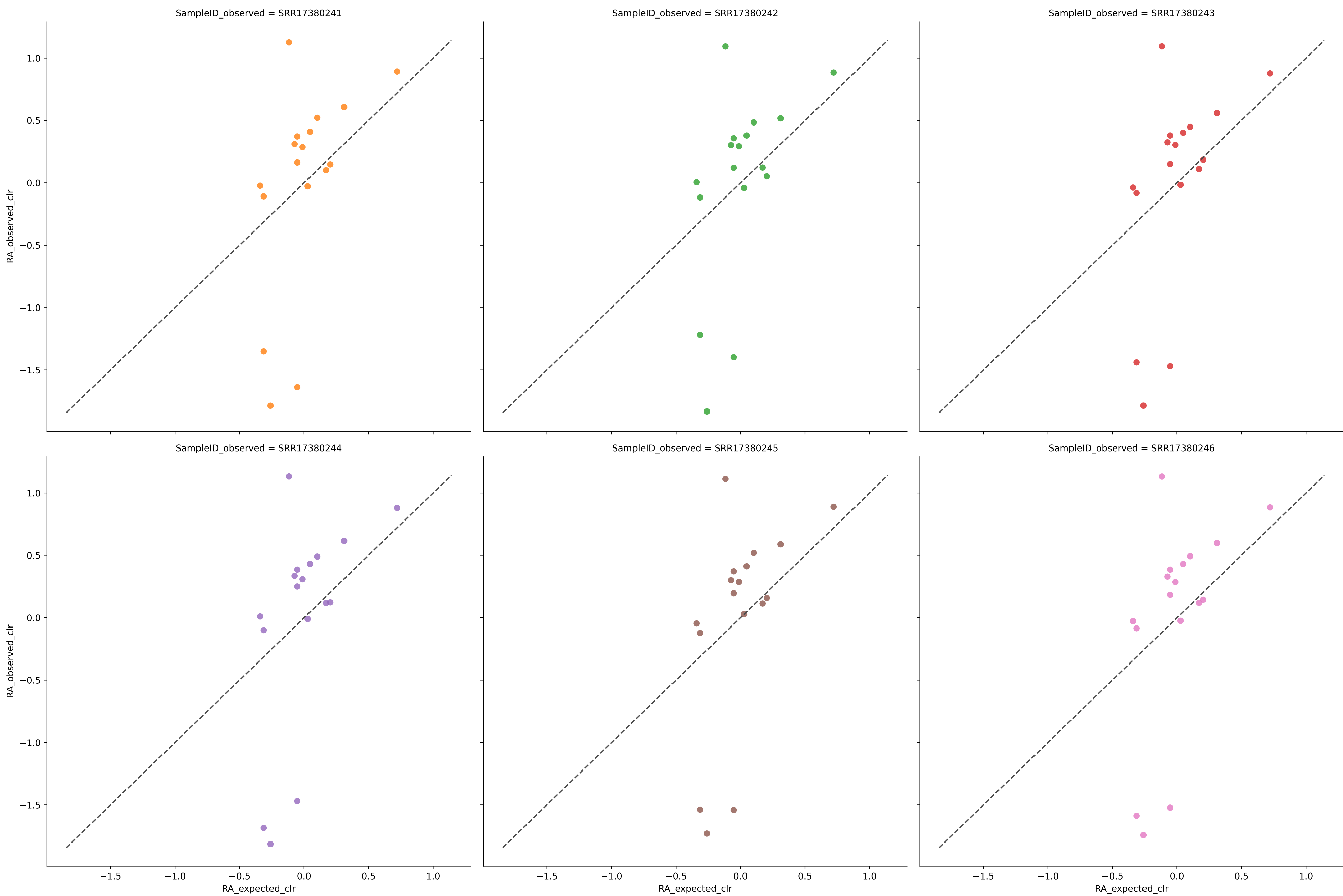


Bivariate Linear Regression for Sample MIX-D in Experiment nist (Species at filter threshold 0.0001)



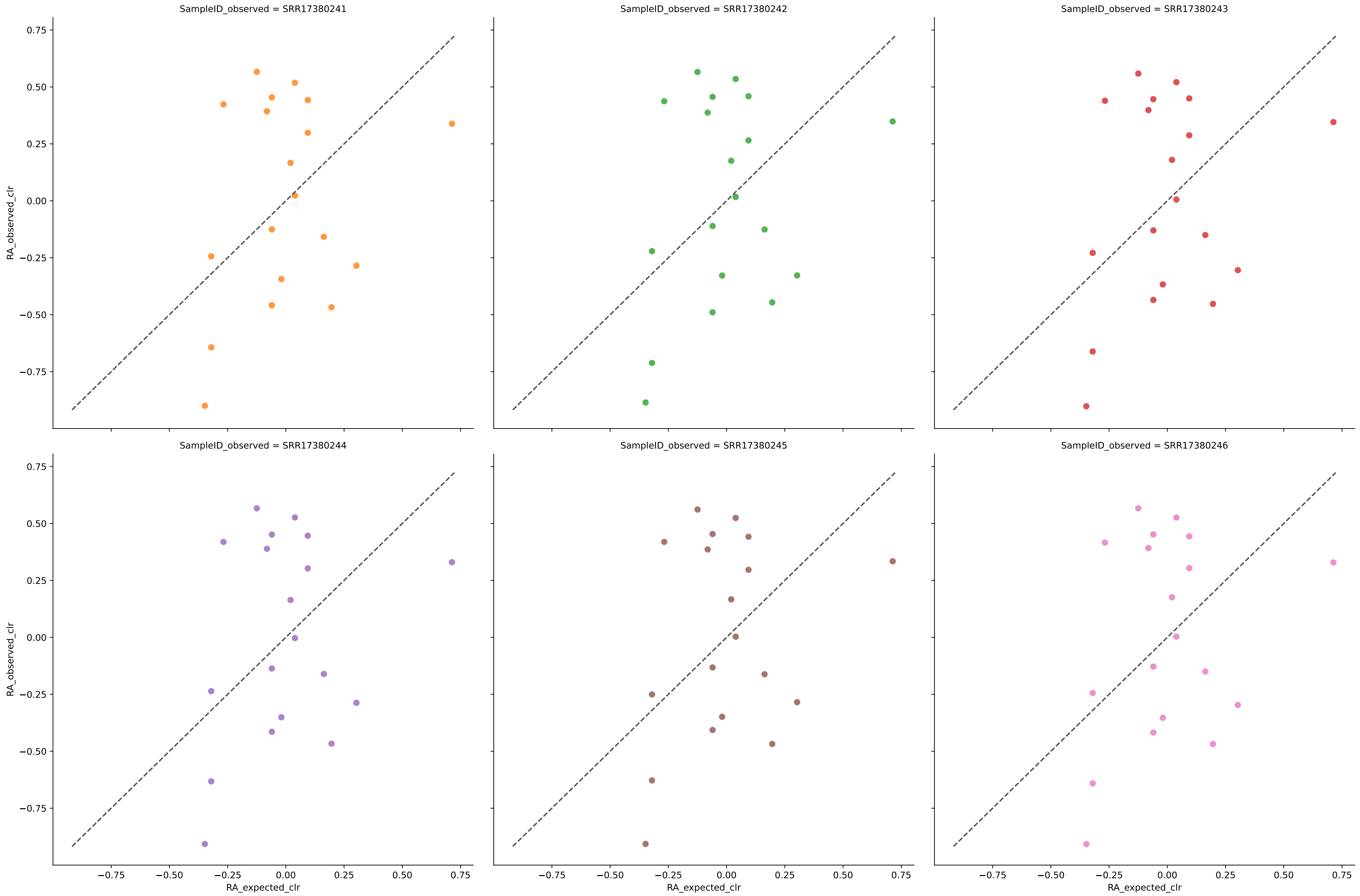
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	7	0.9299	0.0258	1.0255	0.9098	0.0406	100.0000	0.0000
jams	6	0.8432	0.0476	2.4542	0.8573	0.0695	100.0000	0.0000
wgsa2	9	0.8708	0.0325	2.4524	0.8537	0.0584	100.0000	0.0000
woltka	7	0.9911	0.0208	2.9408	0.9272	0.0246	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using bio4 in Experiment tourlousse with filter 0.0001



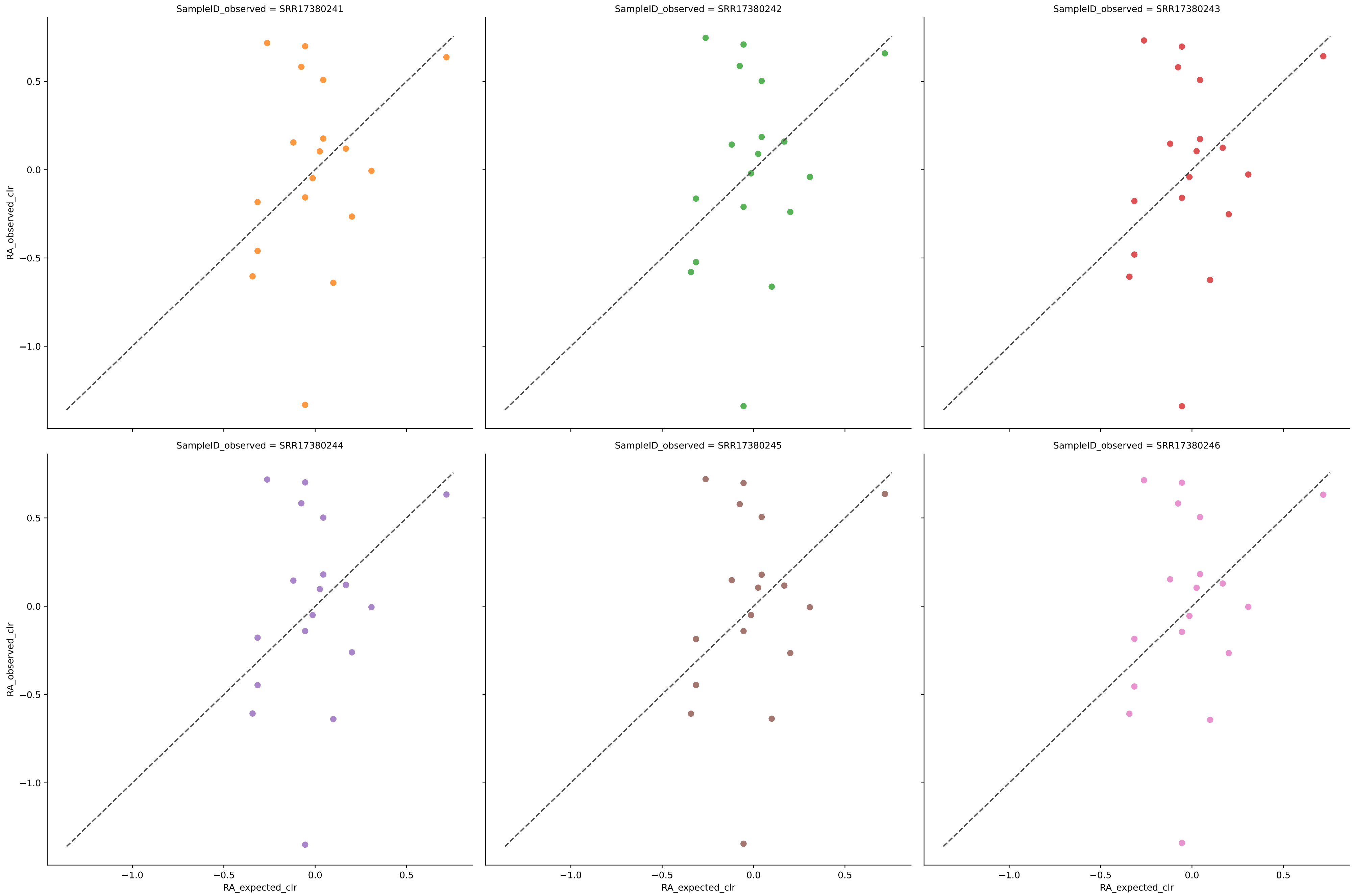
	Diversity	R ²	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR17380241	17	0.2683	0.0031	2.9154	0.8404	0.0048	100.0000	0.0000
SRR17380242	17	0.2667	0.0030	2.7401	0.8446	0.0047	100.0000	0.0000
SRR17380243	17	0.2744	0.0030	2.8350	0.8461	0.0046	100.0000	0.0000
SRR17380244	17	0.2551	0.0032	3.0019	0.8349	0.0048	100.0000	0.0000
SRR17380245	17	0.2771	0.0031	2.8973	0.8429	0.0047	100.0000	0.0000
SRR17380246	17	0.2608	0.0032	2.9331	0.8386	0.0048	100.0000	0.0000
Average	17	0.2671	0.0031	2.8871	0.8412	0.0048	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using jams in Experiment tourlousse with filter 0.0001



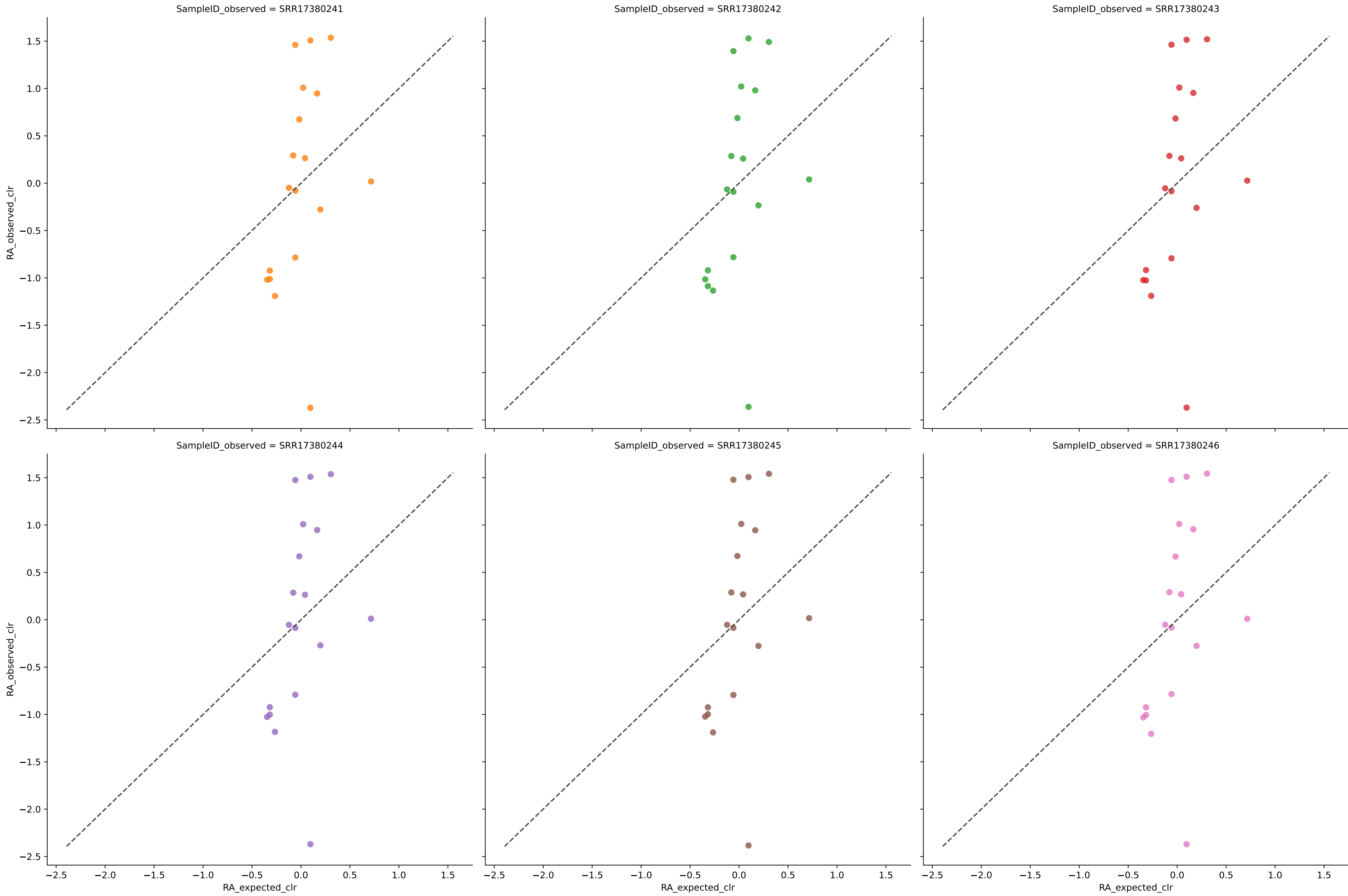
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR17380241	19	0.0294	0.0033	1.8893	0.8104	0.0039	100.0000	0.0000
SRR17380242	19	0.0298	0.0033	1.9110	0.8093	0.0039	100.0000	0.0000
SRR17380243	19	0.0296	0.0033	1.8964	0.8091	0.0039	100.0000	0.0000
SRR17380244	19	0.0276	0.0033	1.8859	0.8093	0.0039	100.0000	0.0000
SRR17380245	19	0.0291	0.0033	1.8776	0.8108	0.0039	100.0000	0.0000
SRR17380246	19	0.0276	0.0033	1.8902	0.8094	0.0039	100.0000	0.0000
Average	19	0.0289	0.0033	1.8917	0.8097	0.0039	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using wgsa in Experiment tourlousse with filter 0.0001



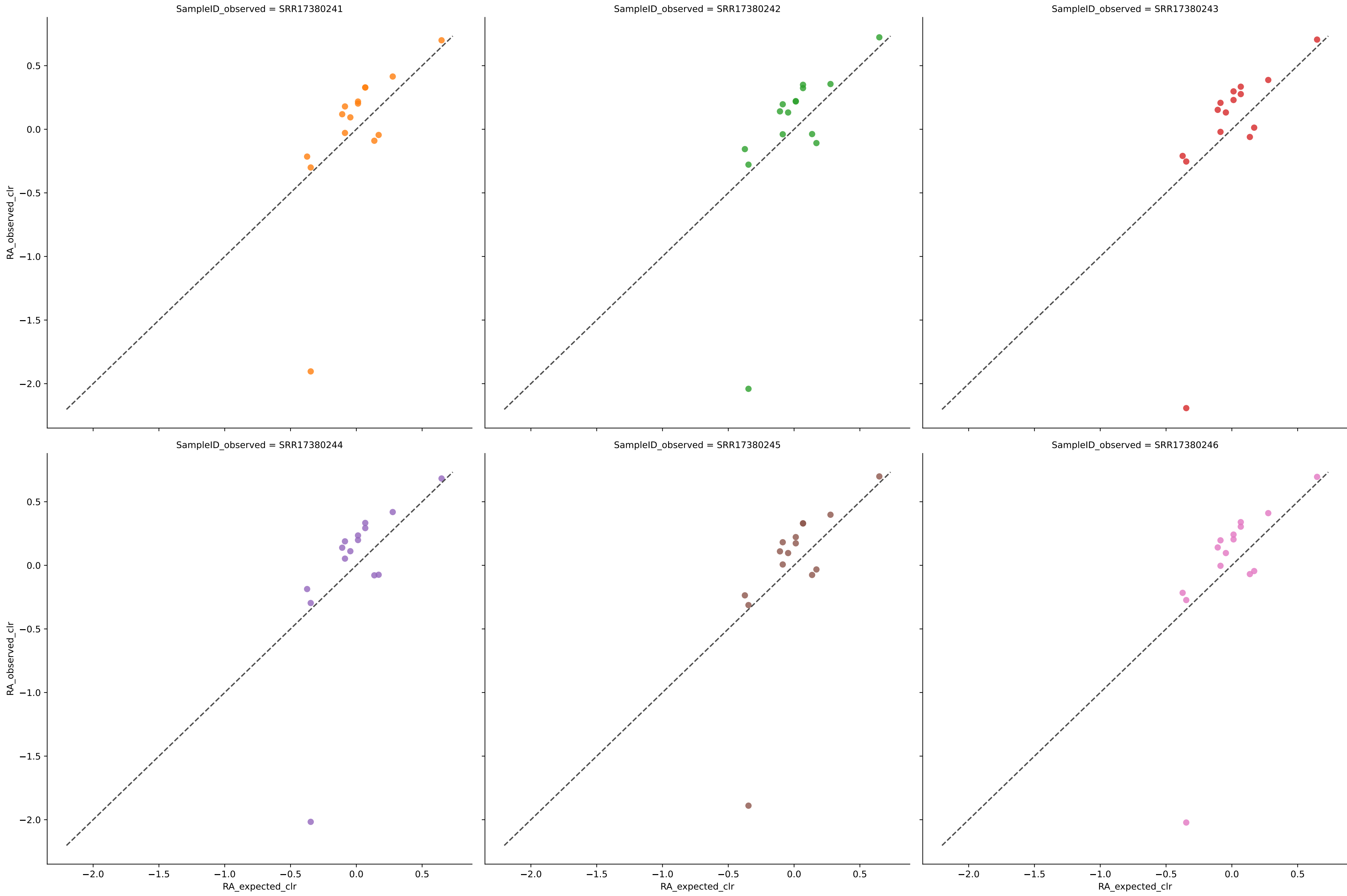
	Diversity	R ²	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR17380241	18	0.0730	0.0034	2.2129	0.8156	0.0044	100.0000	0.0000
SRR17380242	18	0.0747	0.0035	2.2448	0.8130	0.0045	100.0000	0.0000
SRR17380243	18	0.0729	0.0034	2.2182	0.8155	0.0044	100.0000	0.0000
SRR17380244	18	0.0714	0.0034	2.2201	0.8158	0.0044	100.0000	0.0000
SRR17380245	18	0.0727	0.0034	2.2160	0.8165	0.0044	100.0000	0.0000
SRR17380246	18	0.0723	0.0034	2.2157	0.8160	0.0044	100.0000	0.0000
Average	18	0.0728	0.0034	2.2213	0.8154	0.0044	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using woltka in Experiment tourlousse with filter 0.0001



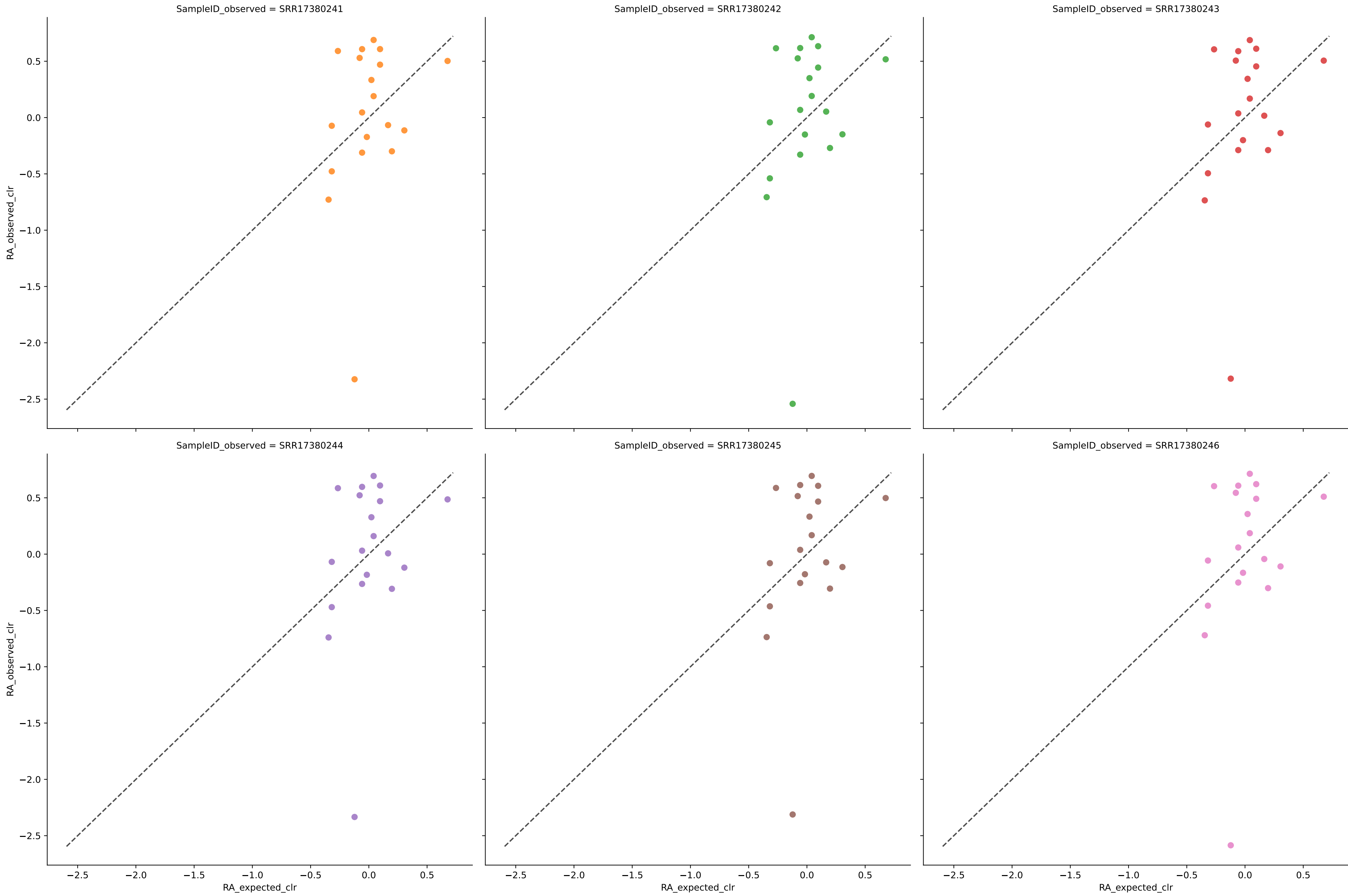
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR17380241	18	0.0619	0.0067	4.1905	0.6387	0.0082	100.0000	0.0000
SRR17380242	18	0.0655	0.0066	4.1572	0.6414	0.0081	100.0000	0.0000
SRR17380243	18	0.0618	0.0067	4.1904	0.6385	0.0082	100.0000	0.0000
SRR17380244	18	0.0608	0.0067	4.1939	0.6374	0.0083	100.0000	0.0000
SRR17380245	18	0.0609	0.0067	4.2048	0.6373	0.0083	100.0000	0.0000
SRR17380246	18	0.0611	0.0067	4.2050	0.6370	0.0083	100.0000	0.0000
Average	18	0.0620	0.0067	4.1903	0.6384	0.0082	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using bio4 in Experiment tourlousse with filter 0.0001



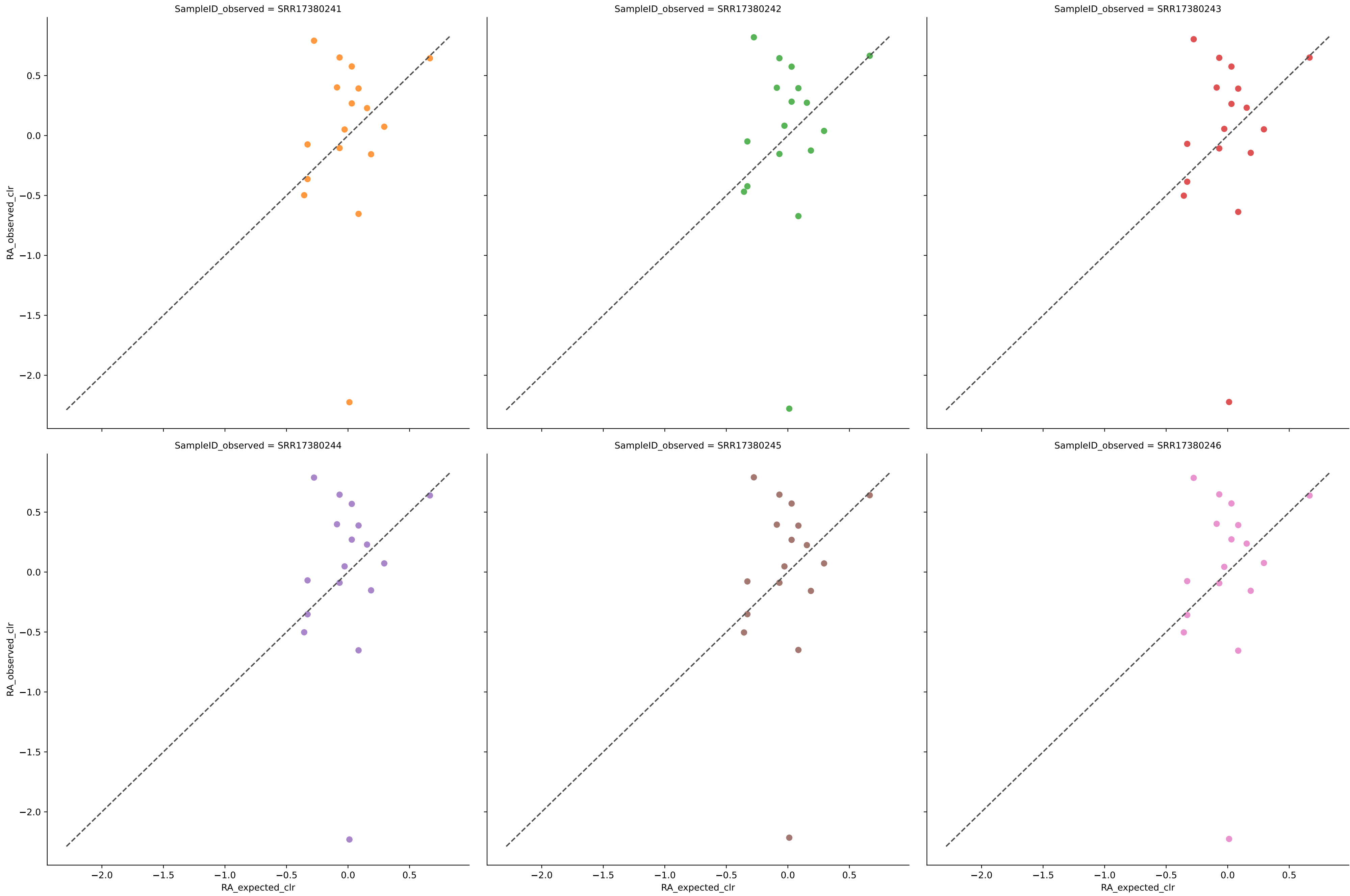
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR17380241	15	0.6588	0.0018	1.7116	0.9196	0.0023	100.0000	0.0000
SRR17380242	15	0.6349	0.0018	1.8567	0.9185	0.0024	100.0000	0.0000
SRR17380243	15	0.6416	0.0018	1.9887	0.9203	0.0024	100.0000	0.0000
SRR17380244	15	0.6284	0.0019	1.8273	0.9153	0.0024	100.0000	0.0000
SRR17380245	15	0.6673	0.0017	1.6908	0.9221	0.0023	100.0000	0.0000
SRR17380246	15	0.6450	0.0018	1.8257	0.9199	0.0024	100.0000	0.0000
Average	15	0.6460	0.0018	1.8168	0.9193	0.0024	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using jams in Experiment tourlousse with filter 0.0001



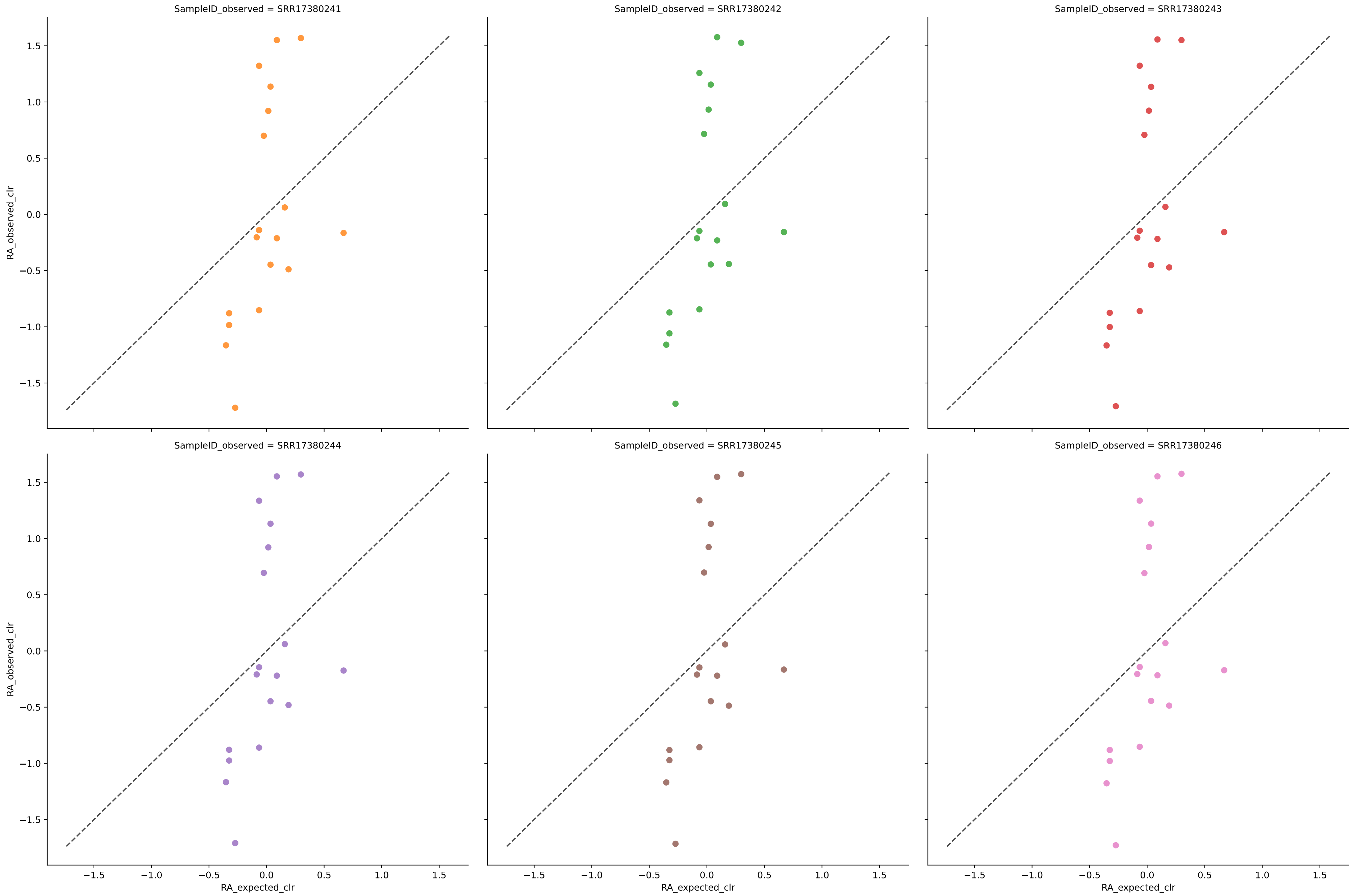
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR17380241	19	0.0685	0.0035	2.8585	0.8028	0.0040	100.0000	0.0000
SRR17380242	19	0.0699	0.0034	3.0443	0.8034	0.0041	100.0000	0.0000
SRR17380243	19	0.0715	0.0034	2.8464	0.8049	0.0040	100.0000	0.0000
SRR17380244	19	0.0671	0.0034	2.8565	0.8044	0.0040	100.0000	0.0000
SRR17380245	19	0.0674	0.0034	2.8435	0.8038	0.0040	100.0000	0.0000
SRR17380246	19	0.0661	0.0035	3.0762	0.8022	0.0041	100.0000	0.0000
Average	19	0.0684	0.0034	2.9209	0.8036	0.0040	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using wgsa in Experiment tourlousse with filter 0.0001



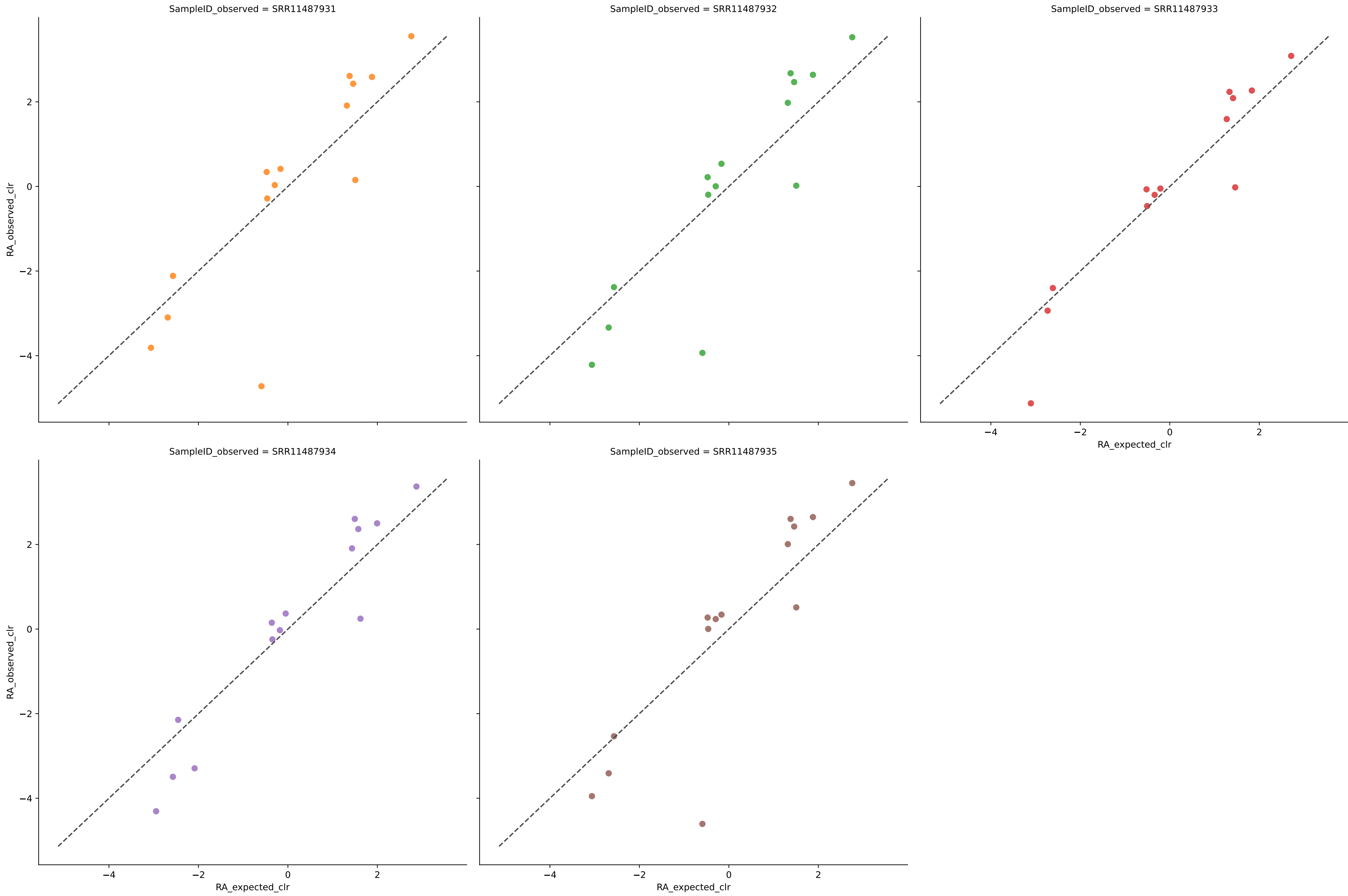
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR17380241	17	0.0588	0.0036	2.8562	0.8157	0.0047	100.0000	0.0000
SRR17380242	17	0.0603	0.0036	2.9180	0.8144	0.0047	100.0000	0.0000
SRR17380243	17	0.0586	0.0036	2.8552	0.8154	0.0047	100.0000	0.0000
SRR17380244	17	0.0579	0.0036	2.8562	0.8166	0.0047	100.0000	0.0000
SRR17380245	17	0.0583	0.0036	2.8440	0.8167	0.0047	100.0000	0.0000
SRR17380246	17	0.0586	0.0036	2.8549	0.8162	0.0047	100.0000	0.0000
Average	17	0.0587	0.0036	2.8641	0.8158	0.0047	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using woltka in Experiment tourlousse with filter 0.0001



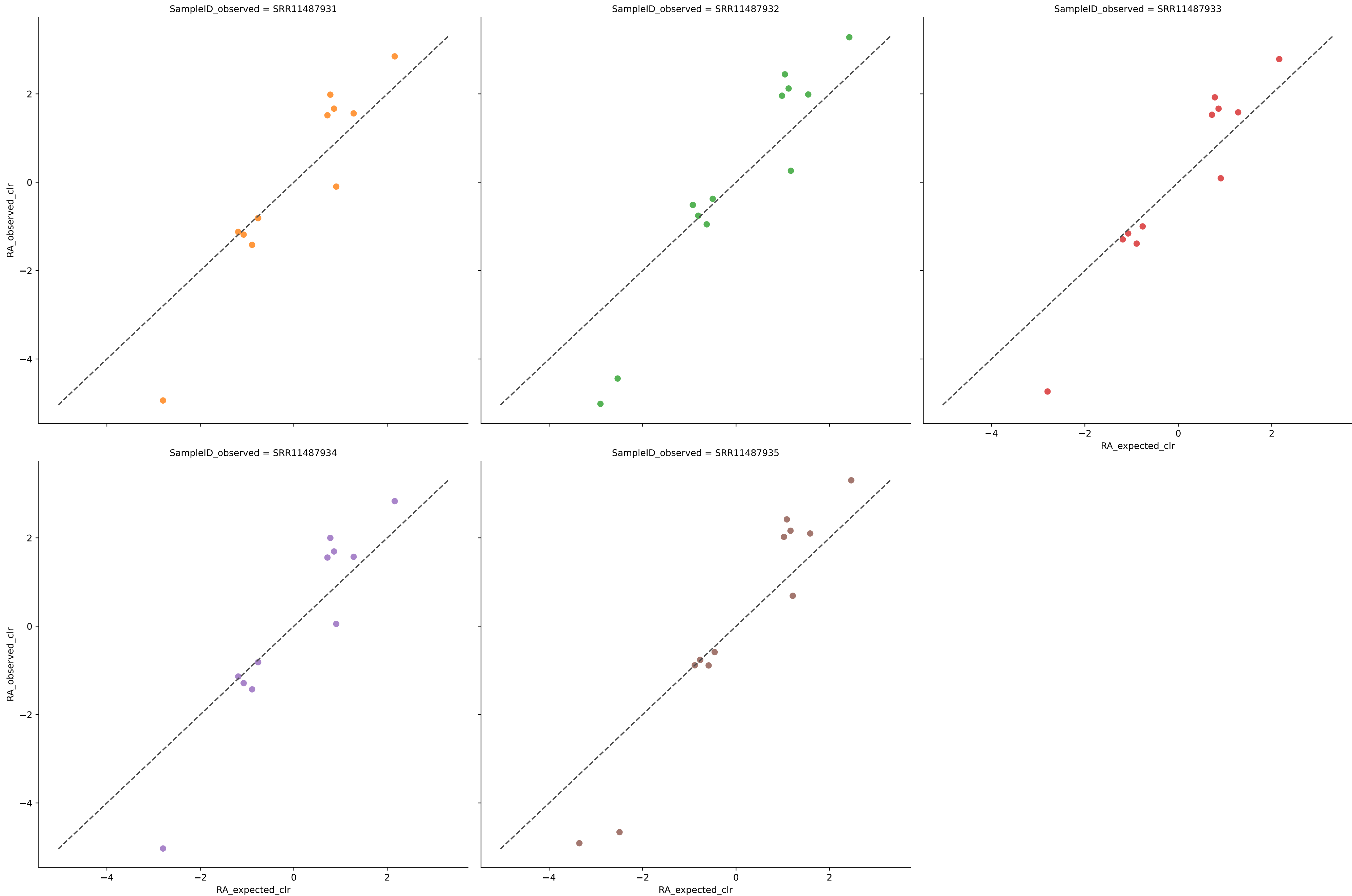
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR17380241	18	0.0560	0.0073	3.7226	0.6054	0.0086	100.0000	0.0000
SRR17380242	18	0.0568	0.0072	3.6942	0.6082	0.0084	100.0000	0.0000
SRR17380243	18	0.0555	0.0073	3.7162	0.6058	0.0086	100.0000	0.0000
SRR17380244	18	0.0550	0.0073	3.7255	0.6044	0.0086	100.0000	0.0000
SRR17380245	18	0.0554	0.0073	3.7281	0.6042	0.0086	100.0000	0.0000
SRR17380246	18	0.0555	0.0073	3.7368	0.6044	0.0086	100.0000	0.0000
Average	18	0.0557	0.0073	3.7206	0.6054	0.0086	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using bio4 in Experiment Amos hilo with filter 0.0001



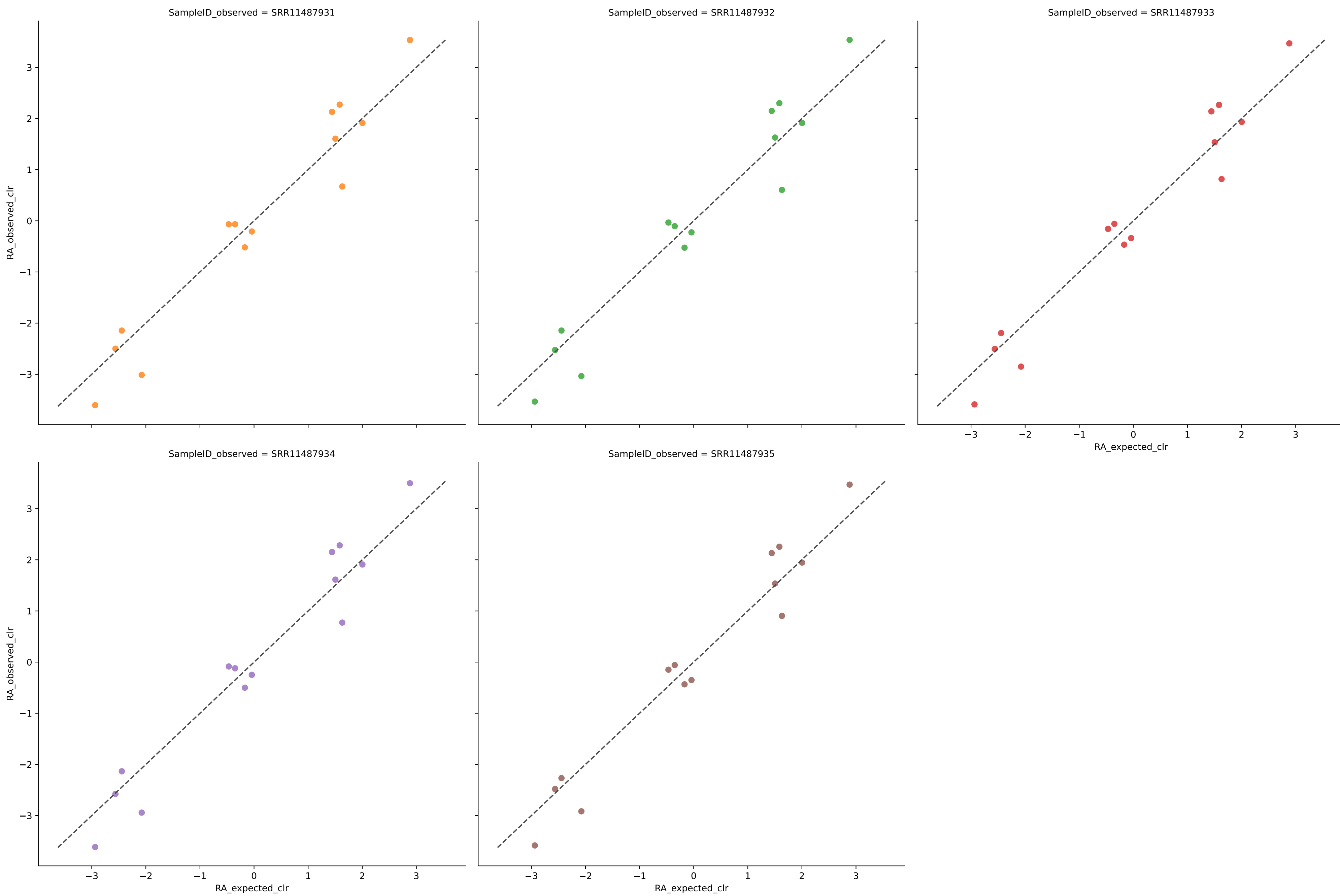
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487931	14	0.9140	0.0037	4.9932	0.8698	0.0066	100.0000	0.0000
SRR11487932	14	0.9020	0.0036	4.5410	0.8729	0.0067	100.0000	0.0000
SRR11487933	13	0.9031	0.0034	2.8846	0.8876	0.0066	100.0000	0.0000
SRR11487934	14	0.8947	0.0032	3.0322	0.8875	0.0067	100.0000	0.0000
SRR11487935	14	0.9105	0.0032	4.8690	0.8887	0.0061	100.0000	0.0000
Average	14	0.9048	0.0034	4.0640	0.8813	0.0065	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using jams in Experiment Amos hilo with filter 0.0001



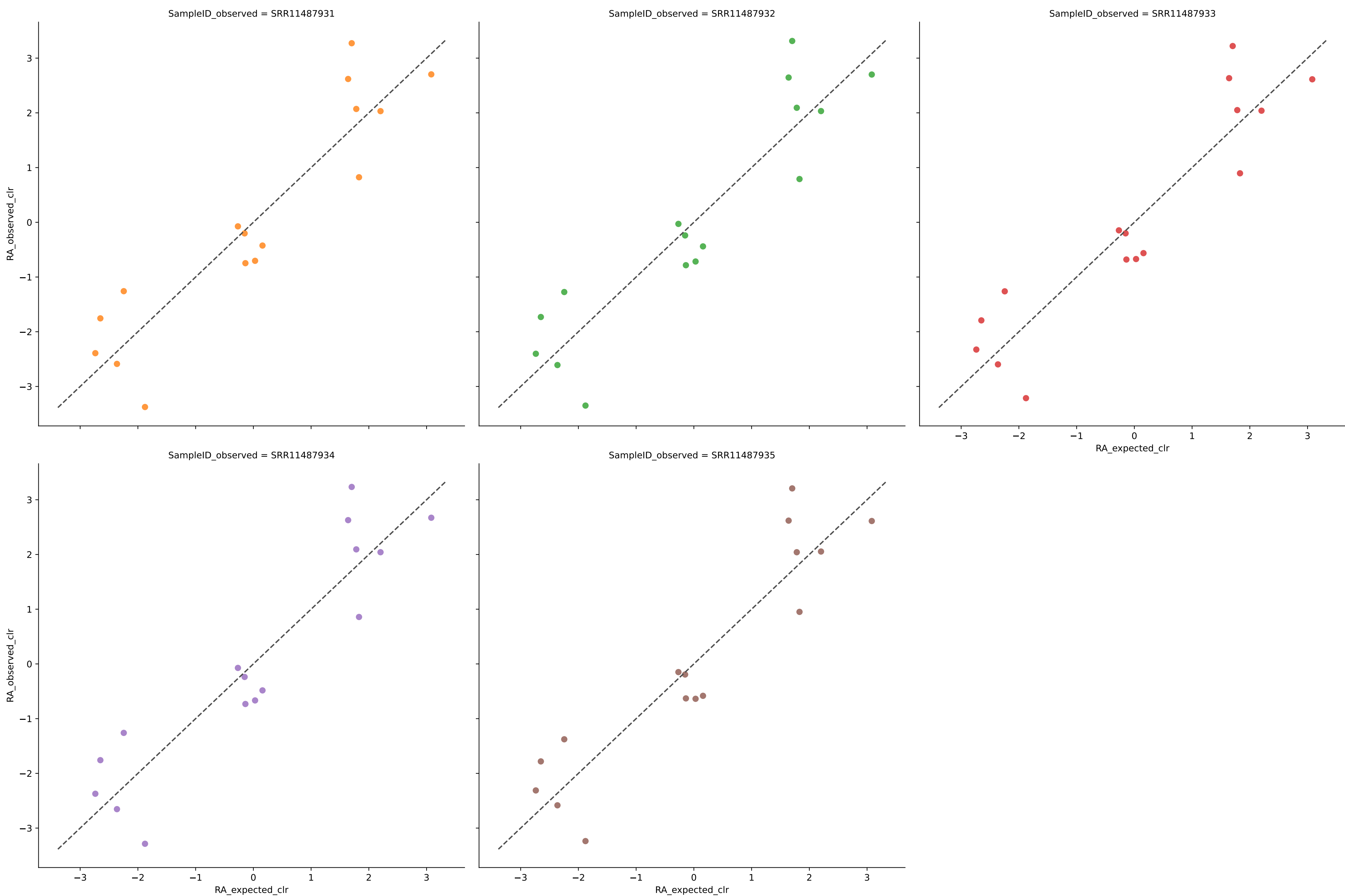
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487931	11	0.8875	0.0060	3.0276	0.8346	0.0082	100.0000	0.0000
SRR11487932	12	0.8852	0.0056	3.7489	0.8320	0.0080	100.0000	0.0000
SRR11487933	11	0.8951	0.0057	2.7982	0.8445	0.0076	100.0000	0.0000
SRR11487934	11	0.8850	0.0059	3.0756	0.8375	0.0081	100.0000	0.0000
SRR11487935	12	0.9068	0.0051	3.4975	0.8457	0.0071	100.0000	0.0000
Average	11	0.8919	0.0057	3.2296	0.8389	0.0078	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using wgsa in Experiment Amos hilo with filter 0.0001



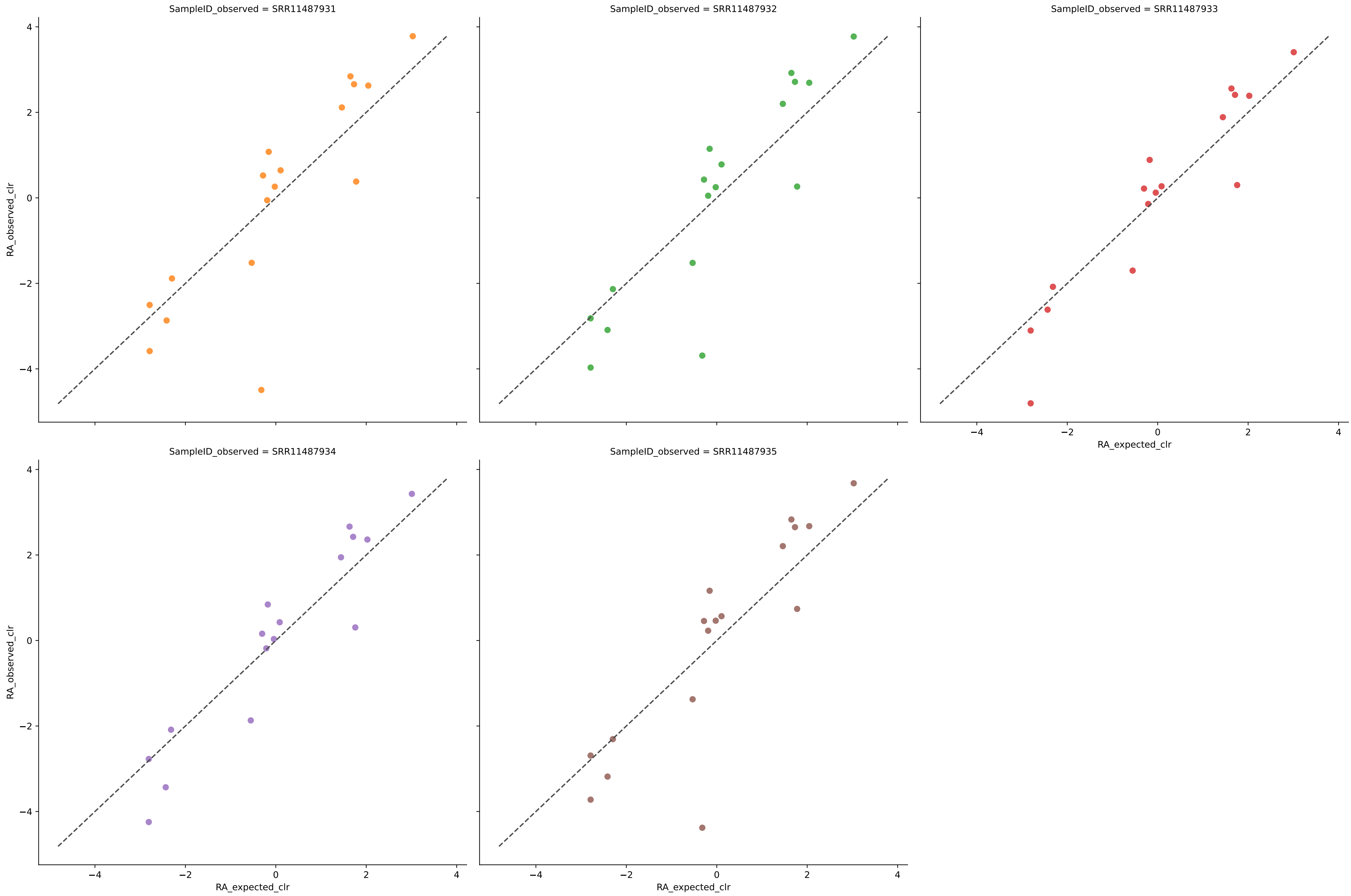
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487931	14	0.9108	0.0053	2.0294	0.8155	0.0087	100.0000	0.0000
SRR11487932	14	0.9085	0.0053	2.0712	0.8139	0.0086	100.0000	0.0000
SRR11487933	14	0.9172	0.0051	1.8476	0.8232	0.0080	100.0000	0.0000
SRR11487934	14	0.9138	0.0051	1.9451	0.8209	0.0082	100.0000	0.0000
SRR11487935	14	0.9210	0.0049	1.8227	0.8270	0.0078	100.0000	0.0000
Average	14	0.9143	0.0051	1.9432	0.8201	0.0083	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using wol in Experiment Amos hilo with filter 0.0001



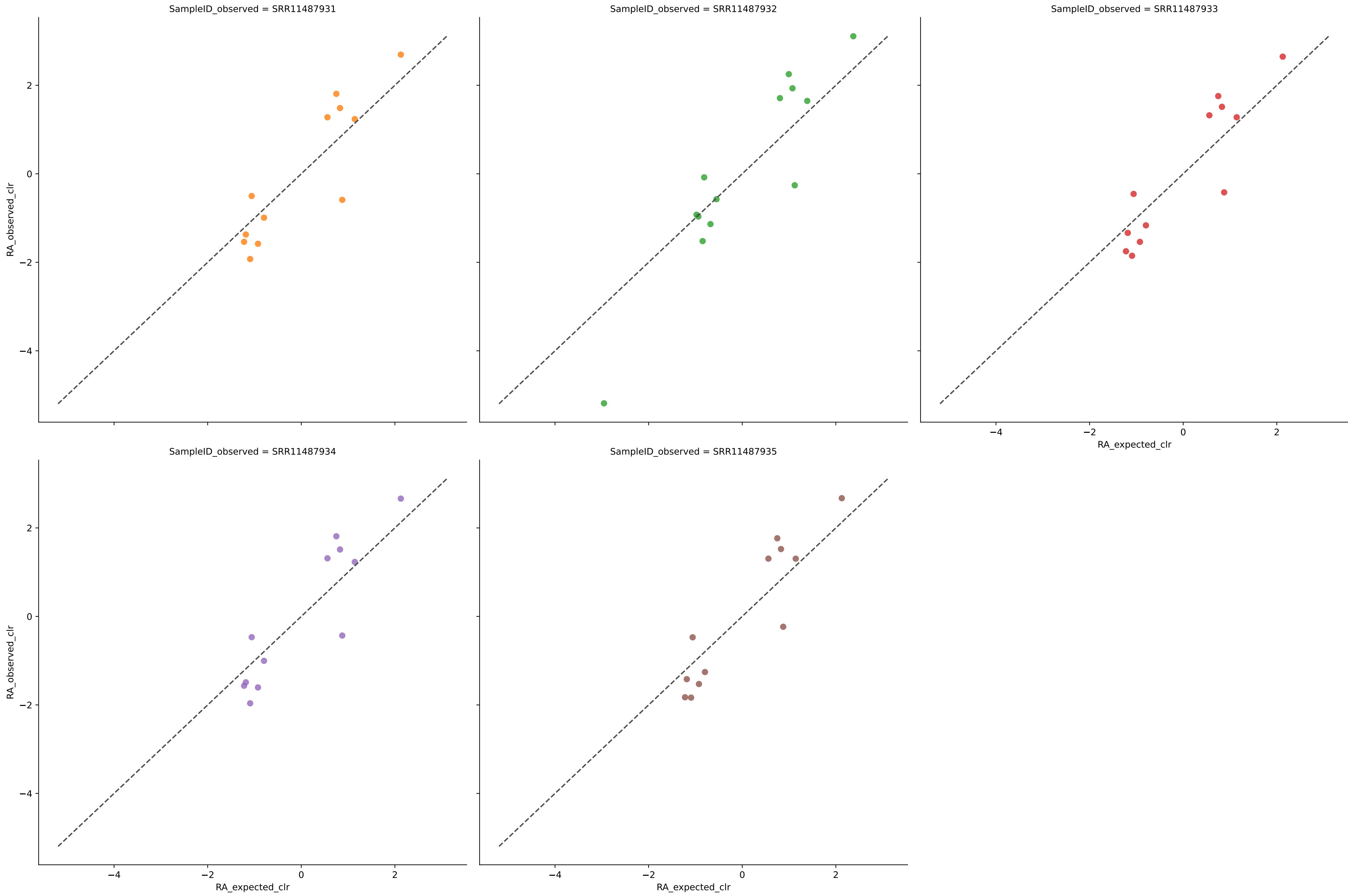
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487931	16	0.3723	0.0086	3.1882	0.6543	0.0166	100.0000	0.0000
SRR11487932	16	0.3532	0.0088	3.2368	0.6454	0.0171	100.0000	0.0000
SRR11487933	16	0.3586	0.0087	3.0849	0.6509	0.0168	100.0000	0.0000
SRR11487934	16	0.3760	0.0086	3.1331	0.6548	0.0165	100.0000	0.0000
SRR11487935	16	0.3660	0.0086	3.0234	0.6564	0.0166	100.0000	0.0000
Average	16	0.3652	0.0087	3.1333	0.6524	0.0167	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using bio4 in Experiment Amos hilo with filter 0.0001



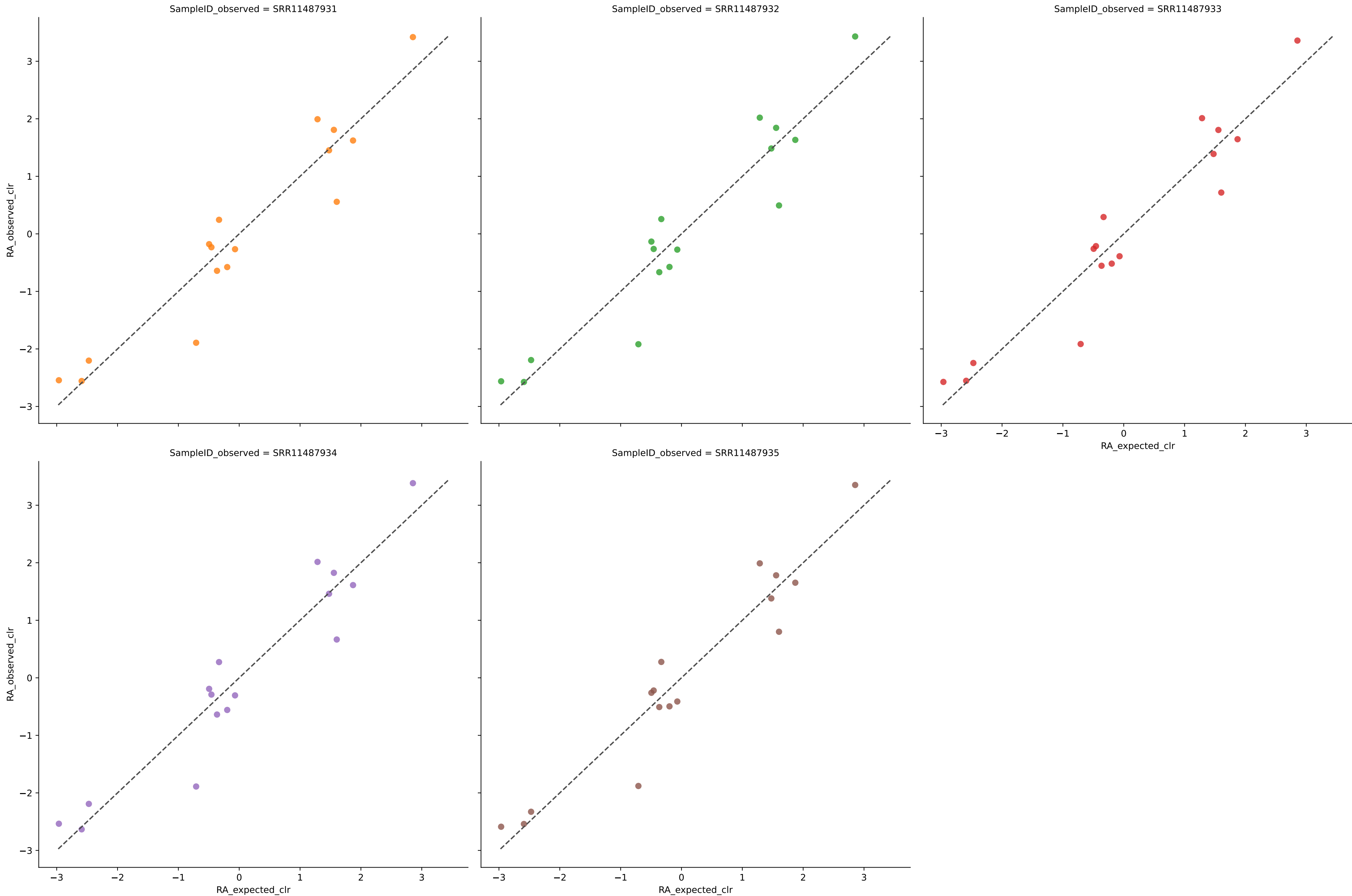
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487931	17	0.9142	0.0034	5.2528	0.8578	0.0060	100.0000	0.0000
SRR11487932	17	0.9028	0.0032	4.8354	0.8626	0.0061	100.0000	0.0000
SRR11487933	16	0.9054	0.0030	3.3029	0.8786	0.0060	100.0000	0.0000
SRR11487934	16	0.8930	0.0032	3.2327	0.8697	0.0063	100.0000	0.0000
SRR11487935	17	0.9113	0.0029	5.1244	0.8774	0.0056	100.0000	0.0000
Average	17	0.9054	0.0032	4.3496	0.8692	0.0060	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using jams in Experiment Amos hilo with filter 0.0001



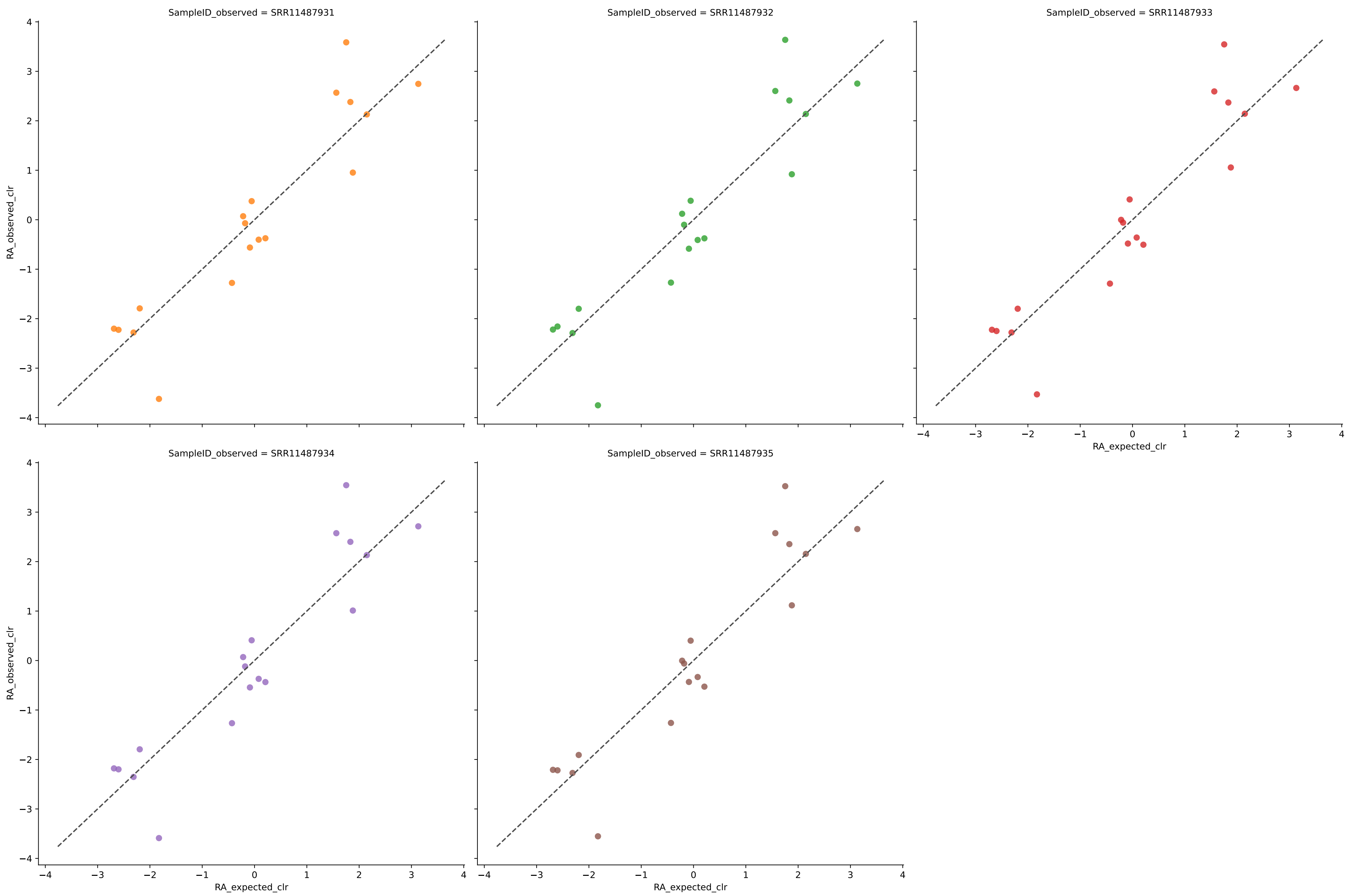
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487931	12	0.8855	0.0060	2.4794	0.8201	0.0083	100.0000	0.0000
SRR11487932	13	0.8822	0.0056	3.4408	0.8168	0.0081	100.0000	0.0000
SRR11487933	12	0.8897	0.0058	2.4059	0.8272	0.0078	100.0000	0.0000
SRR11487934	12	0.8812	0.0059	2.4483	0.8215	0.0081	100.0000	0.0000
SRR11487935	12	0.8978	0.0057	2.3490	0.8301	0.0076	100.0000	0.0000
Average	12	0.8873	0.0058	2.6247	0.8231	0.0080	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using wgsa in Experiment Amos hilo with filter 0.0001



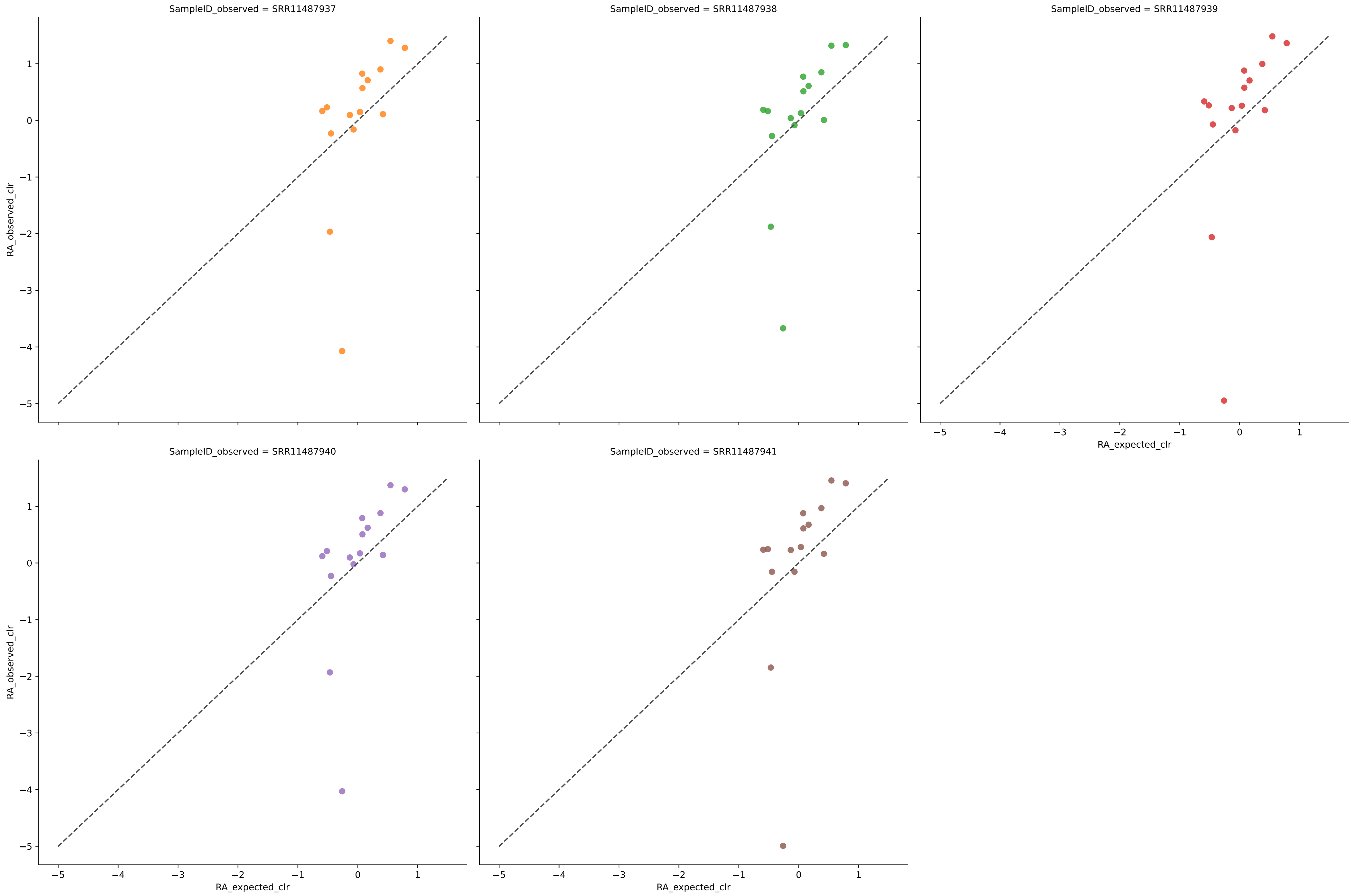
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487931	16	0.9115	0.0047	2.1050	0.8138	0.0087	100.0000	0.0000
SRR11487932	16	0.9101	0.0047	2.1750	0.8125	0.0086	100.0000	0.0000
SRR11487933	16	0.9178	0.0044	2.0224	0.8225	0.0079	100.0000	0.0000
SRR11487934	16	0.9149	0.0045	2.0573	0.8187	0.0082	100.0000	0.0000
SRR11487935	16	0.9215	0.0043	1.9332	0.8271	0.0078	100.0000	0.0000
Average	16	0.9152	0.0045	2.0586	0.8189	0.0082	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using wol in Experiment Amos hilo with filter 0.0001



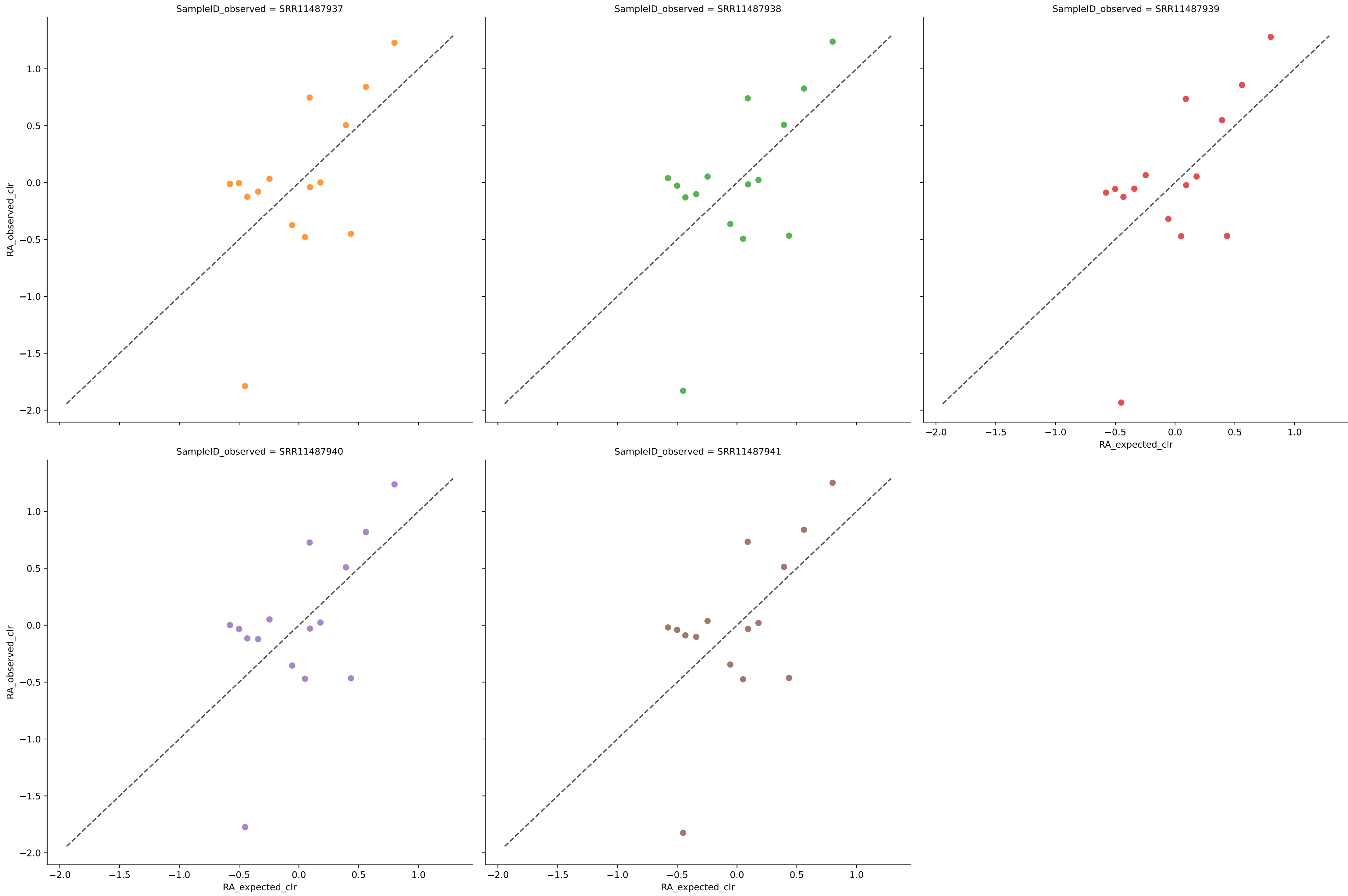
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487931	18	0.2914	0.0083	3.3507	0.6248	0.0177	100.0000	0.0000
SRR11487932	18	0.2756	0.0085	3.4846	0.6160	0.0181	100.0000	0.0000
SRR11487933	18	0.2816	0.0084	3.2745	0.6226	0.0177	100.0000	0.0000
SRR11487934	18	0.2963	0.0083	3.3150	0.6261	0.0175	100.0000	0.0000
SRR11487935	18	0.2883	0.0083	3.2343	0.6285	0.0176	100.0000	0.0000
Average	18	0.2866	0.0084	3.3318	0.6236	0.0177	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using bio4 in Experiment Amos mixed with filter 0.0001



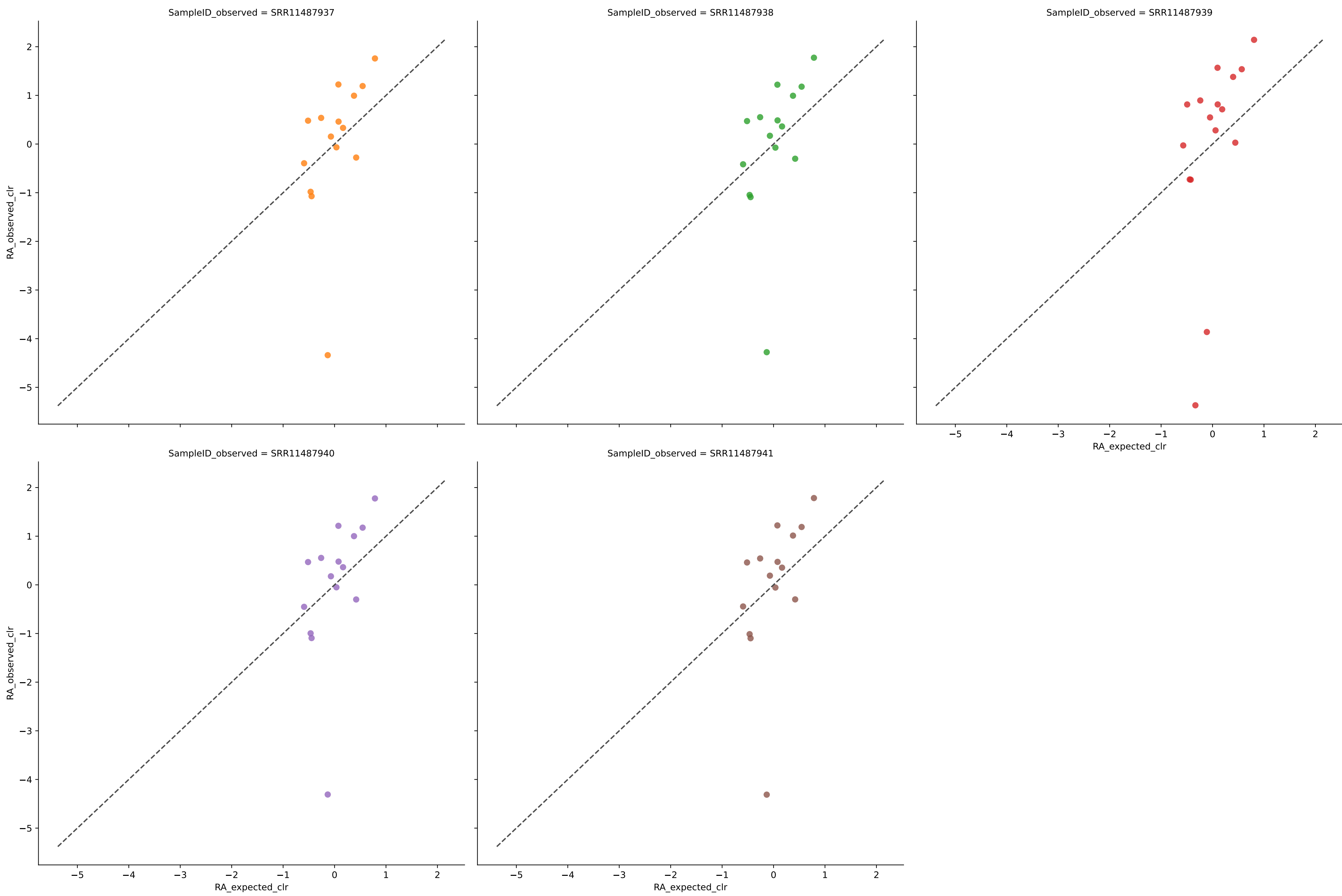
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487937	15	0.6595	0.0047	4.5249	0.8236	0.0058	100.0000	0.0000
SRR11487938	15	0.6856	0.0046	4.1118	0.8260	0.0056	100.0000	0.0000
SRR11487939	15	0.6543	0.0046	5.3975	0.8269	0.0058	100.0000	0.0000
SRR11487940	15	0.6903	0.0044	4.4374	0.8352	0.0055	100.0000	0.0000
SRR11487941	15	0.6856	0.0046	5.3530	0.8283	0.0057	100.0000	0.0000
Average	15	0.6751	0.0046	4.7649	0.8280	0.0057	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using jams in Experiment Amos mixed with filter 0.0001



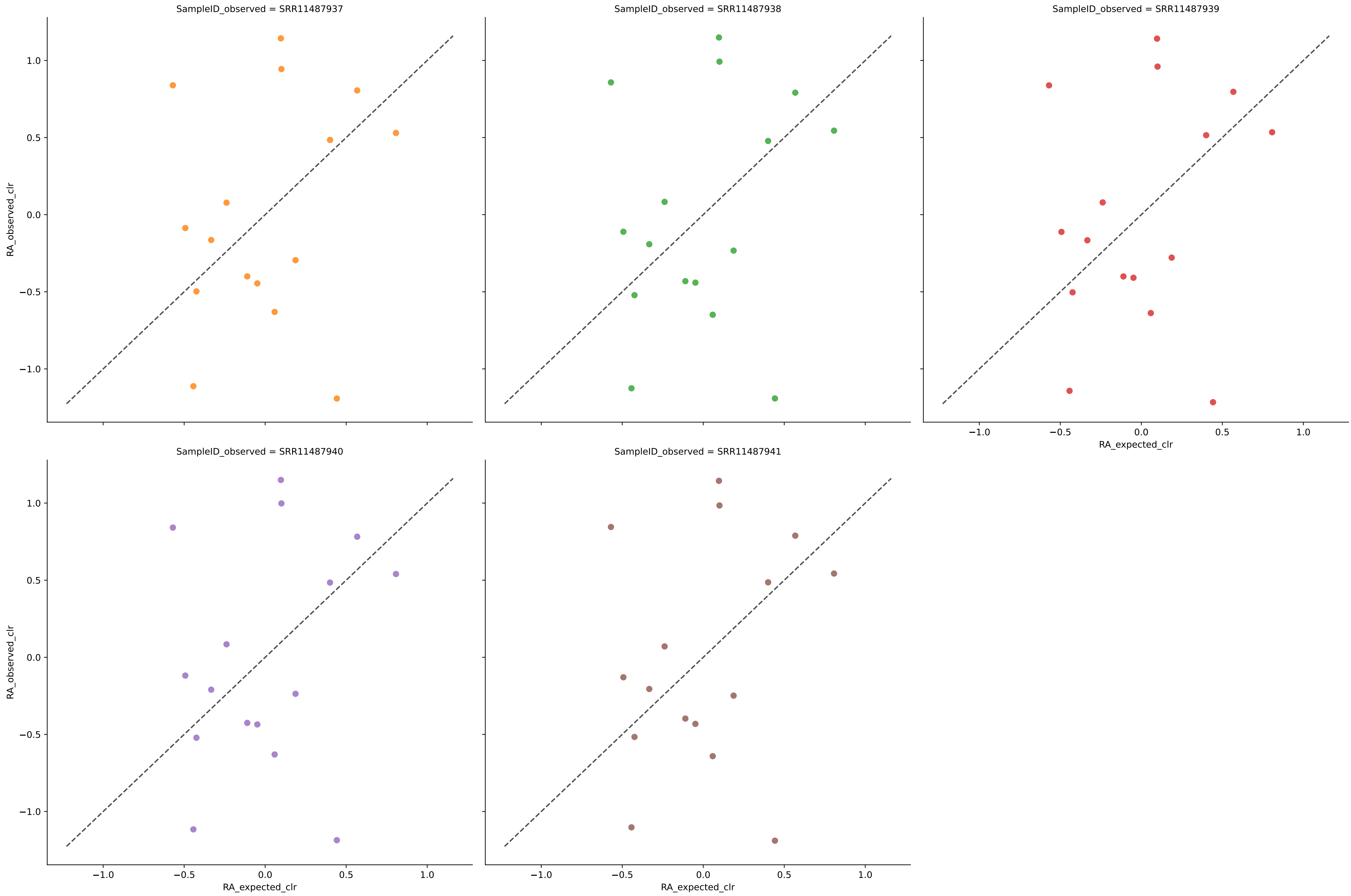
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487937	15	0.5720	0.0047	2.1252	0.8246	0.0057	100.0000	0.0000
SRR11487938	15	0.5697	0.0047	2.1639	0.8250	0.0058	100.0000	0.0000
SRR11487939	15	0.5984	0.0046	2.2036	0.8278	0.0057	100.0000	0.0000
SRR11487940	15	0.5777	0.0046	2.1051	0.8269	0.0057	100.0000	0.0000
SRR11487941	15	0.5820	0.0046	2.1406	0.8259	0.0058	100.0000	0.0000
Average	15	0.5800	0.0046	2.1477	0.8260	0.0058	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using wgsa in Experiment Amos mixed with filter 0.0001



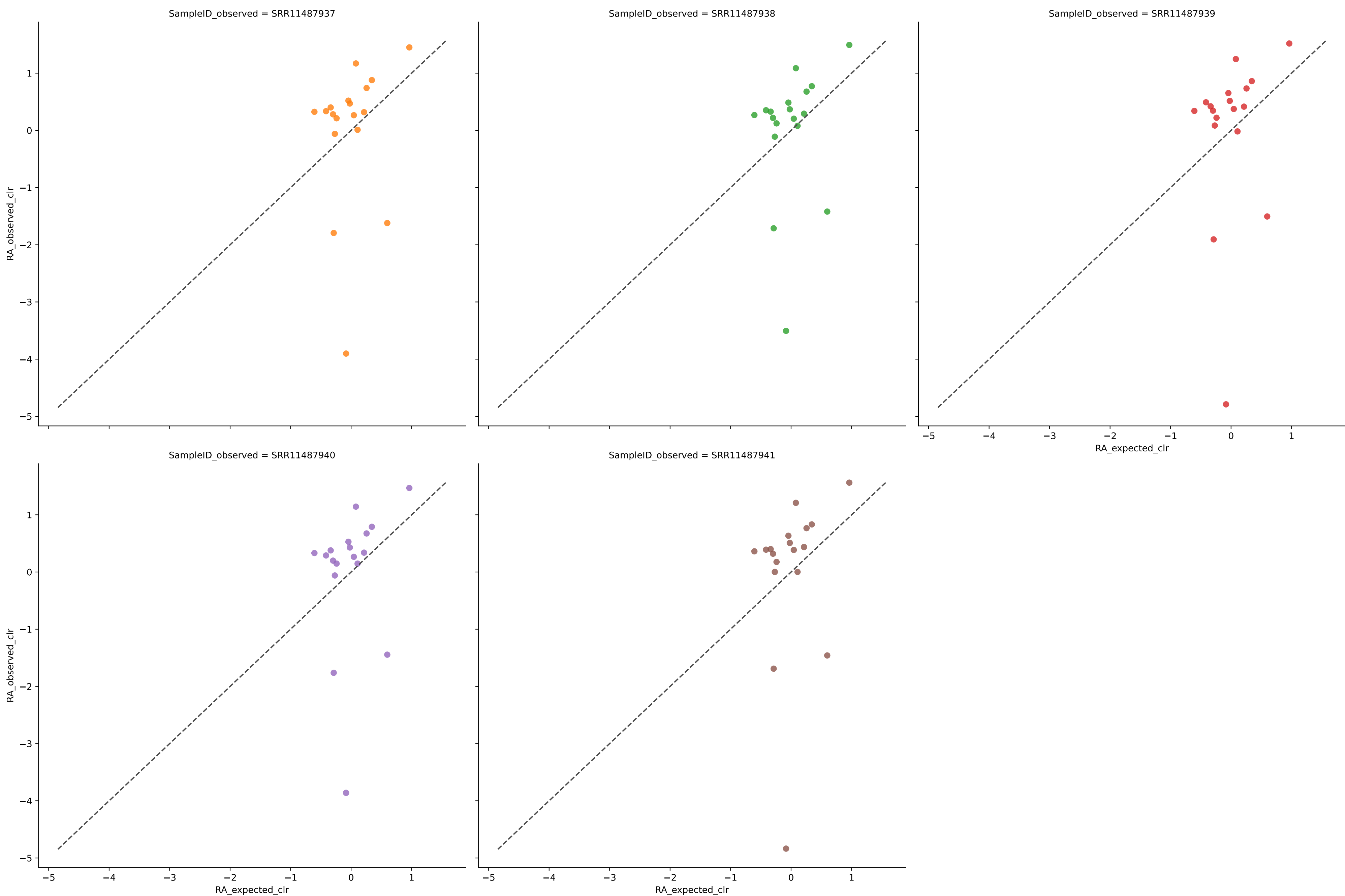
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487937	15	0.5991	0.0063	4.8794	0.7611	0.0080	100.0000	0.0000
SRR11487938	15	0.6002	0.0063	4.8442	0.7613	0.0081	100.0000	0.0000
SRR11487939	16	0.6217	0.0065	7.0792	0.7436	0.0081	100.0000	0.0000
SRR11487940	15	0.6031	0.0063	4.8625	0.7618	0.0080	100.0000	0.0000
SRR11487941	15	0.6063	0.0064	4.8704	0.7602	0.0081	100.0000	0.0000
Average	15	0.6061	0.0064	5.3072	0.7576	0.0081	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using wol in Experiment Amos mixed with filter 0.0001



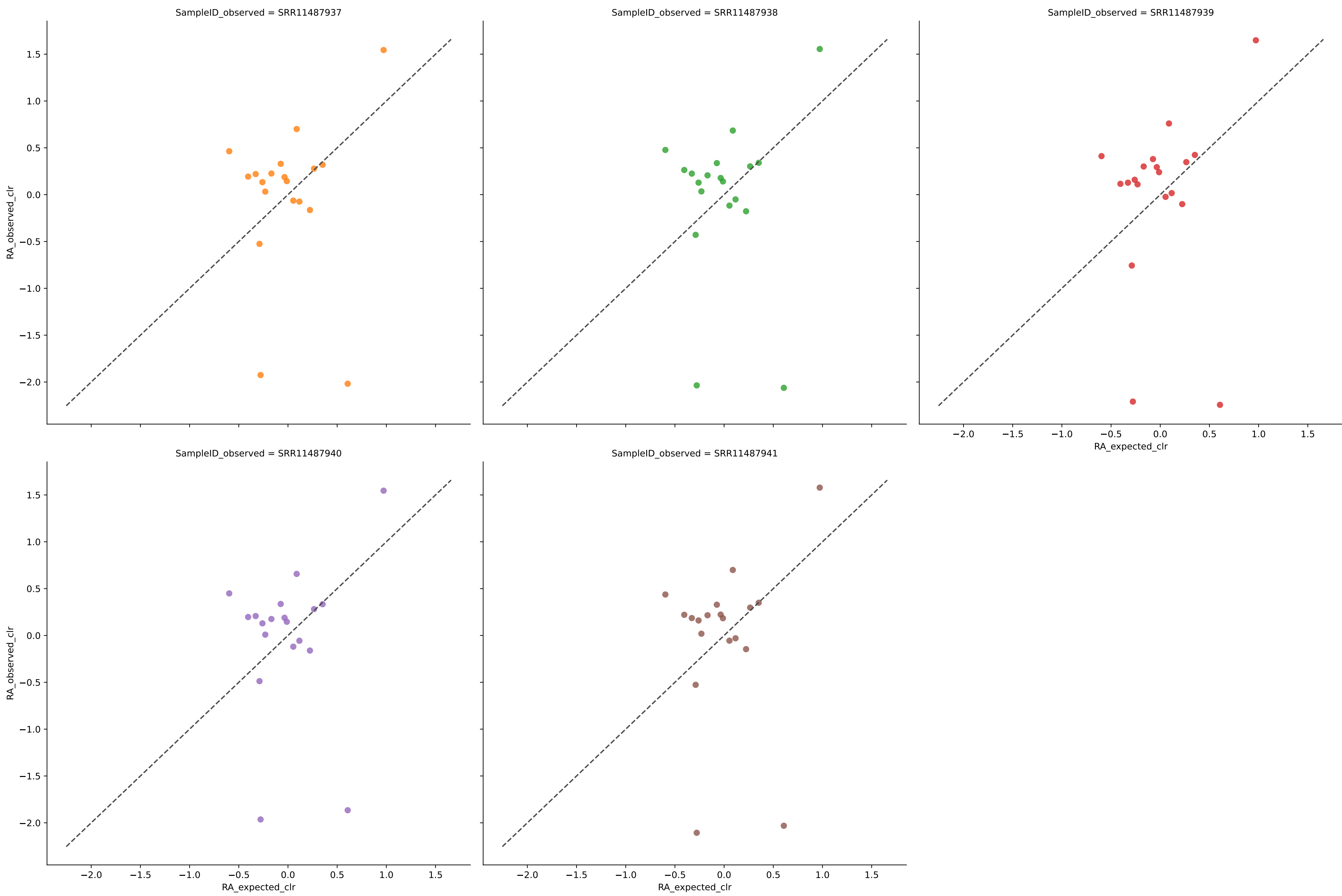
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487937	16	0.0645	0.0066	2.8809	0.7364	0.0087	100.0000	0.0000
SRR11487938	16	0.0633	0.0066	2.9006	0.7340	0.0089	100.0000	0.0000
SRR11487939	16	0.0662	0.0065	2.8965	0.7387	0.0087	100.0000	0.0000
SRR11487940	16	0.0643	0.0066	2.8819	0.7351	0.0088	100.0000	0.0000
SRR11487941	16	0.0658	0.0066	2.8748	0.7369	0.0088	100.0000	0.0000
Average	16	0.0648	0.0066	2.8869	0.7362	0.0088	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using bio4 in Experiment Amos mixed with filter 0.0001



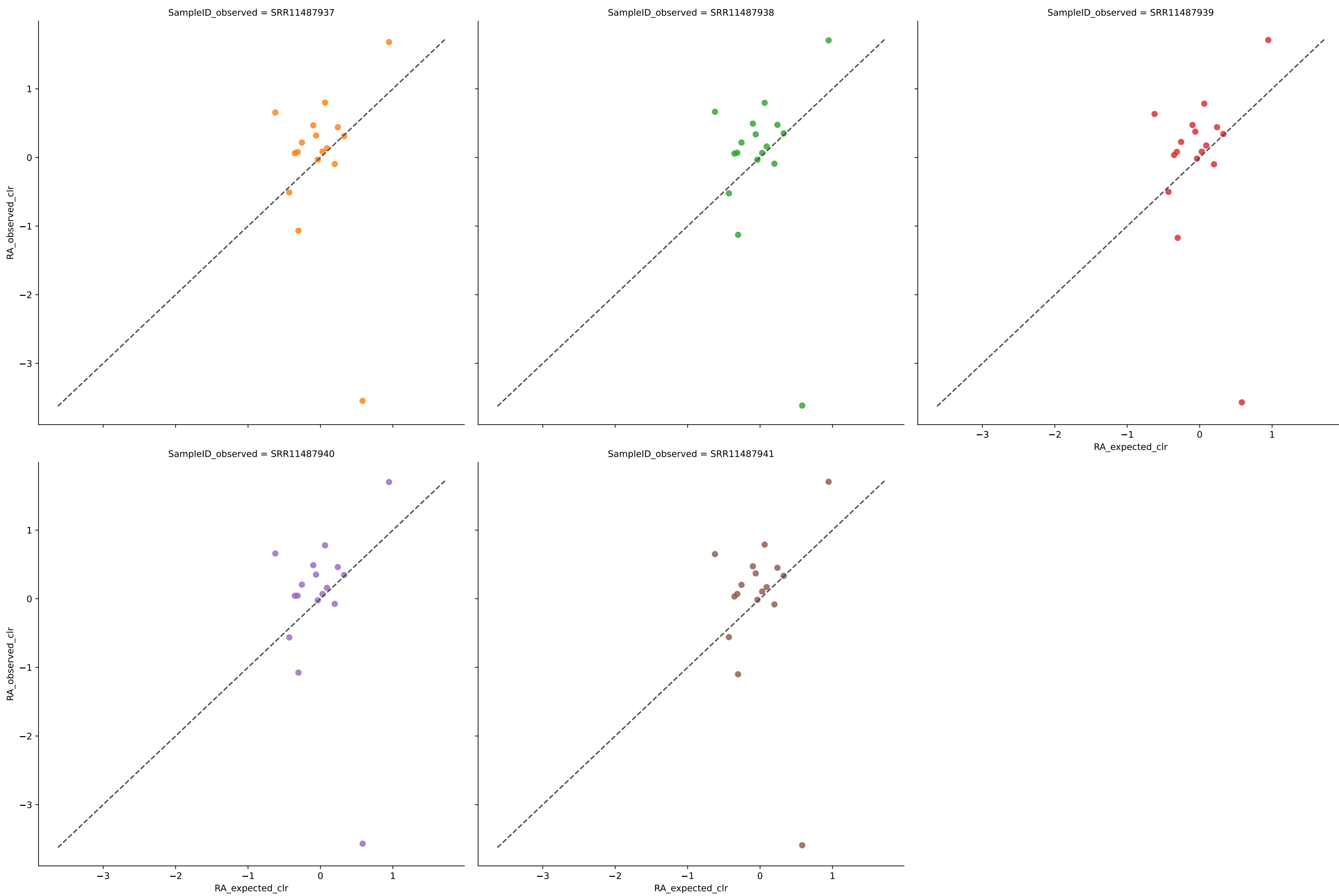
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487937	18	0.2996	0.0046	5.1875	0.7914	0.0063	100.0000	0.0000
SRR11487938	18	0.3547	0.0045	4.7066	0.7969	0.0062	100.0000	0.0000
SRR11487939	18	0.2928	0.0046	5.9450	0.7951	0.0063	100.0000	0.0000
SRR11487940	18	0.3277	0.0044	5.0307	0.8030	0.0061	100.0000	0.0000
SRR11487941	18	0.3332	0.0045	5.8823	0.7971	0.0063	100.0000	0.0000
Average	18	0.3216	0.0045	5.3504	0.7967	0.0062	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using jams in Experiment Amos mixed with filter 0.0001



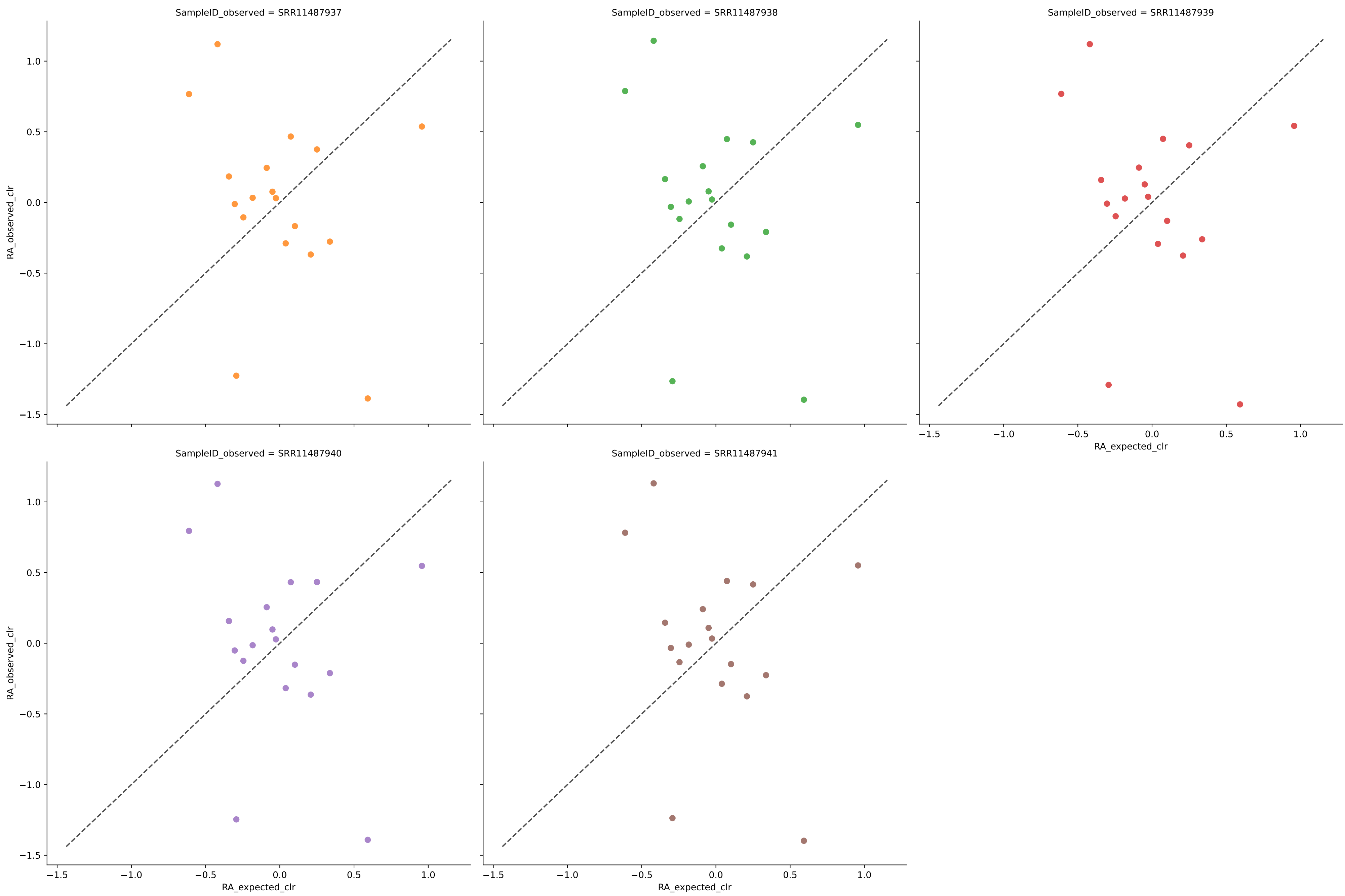
	Diversity	R ²	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487937	19	0.3708	0.0043	3.6003	0.7947	0.0060	100.0000	0.0000
SRR11487938	19	0.3682	0.0044	3.6950	0.7941	0.0061	100.0000	0.0000
SRR11487939	19	0.4162	0.0042	3.9369	0.7974	0.0060	100.0000	0.0000
SRR11487940	19	0.3834	0.0043	3.4885	0.7968	0.0060	100.0000	0.0000
SRR11487941	19	0.3887	0.0043	3.6944	0.7964	0.0060	100.0000	0.0000
Average	19	0.3854	0.0043	3.6830	0.7959	0.0060	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using wgsa in Experiment Amos mixed with filter 0.0001



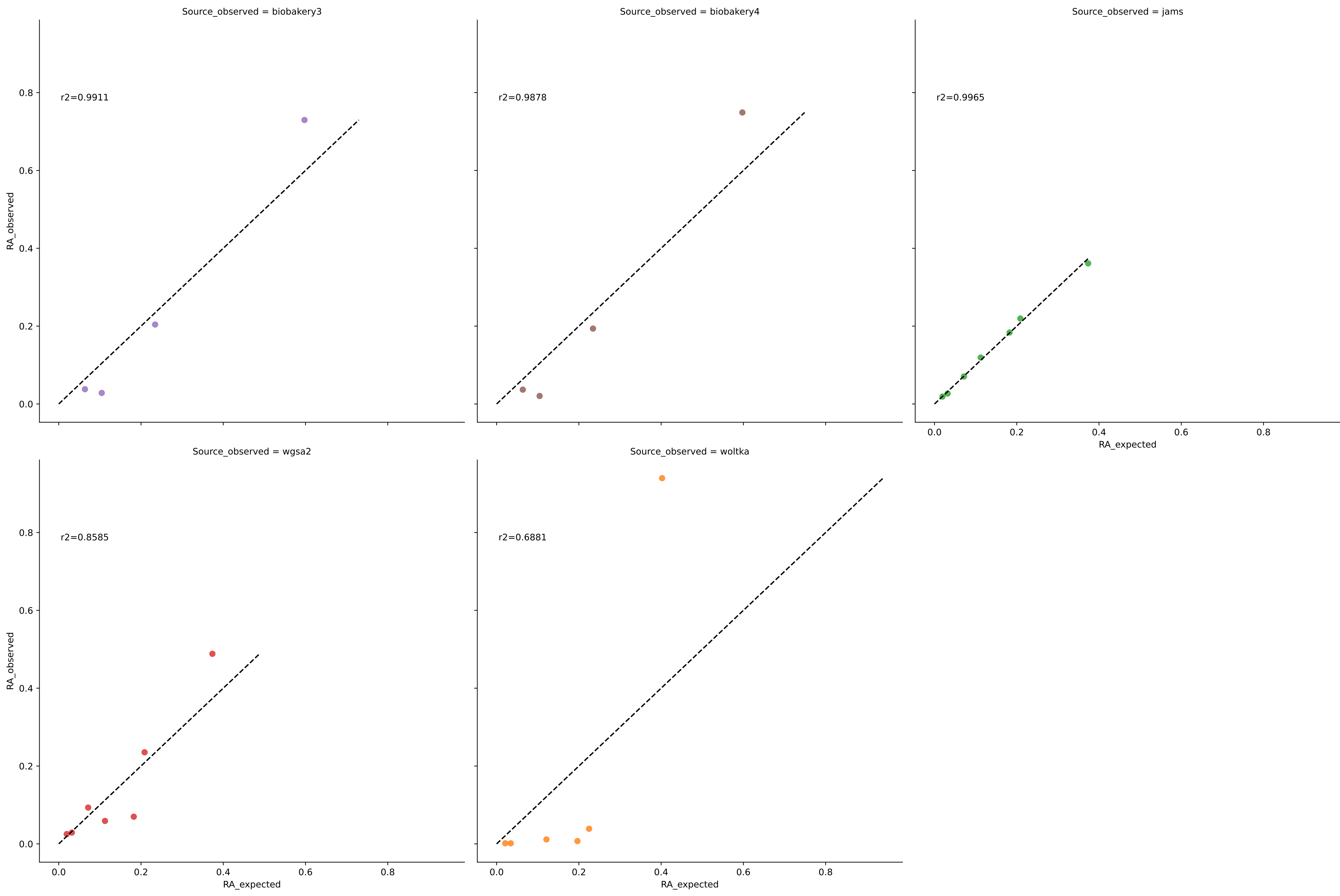
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487937	17	0.3833	0.0051	4.6362	0.7831	0.0074	100.0000	0.0000
SRR11487938	17	0.3924	0.0051	4.7186	0.7829	0.0074	100.0000	0.0000
SRR11487939	17	0.3983	0.0051	4.6751	0.7827	0.0075	100.0000	0.0000
SRR11487940	17	0.3953	0.0051	4.6599	0.7843	0.0074	100.0000	0.0000
SRR11487941	17	0.3961	0.0051	4.6848	0.7841	0.0074	100.0000	0.0000
Average	17	0.3931	0.0051	4.6749	0.7834	0.0074	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using wol in Experiment Amos mixed with filter 0.0001

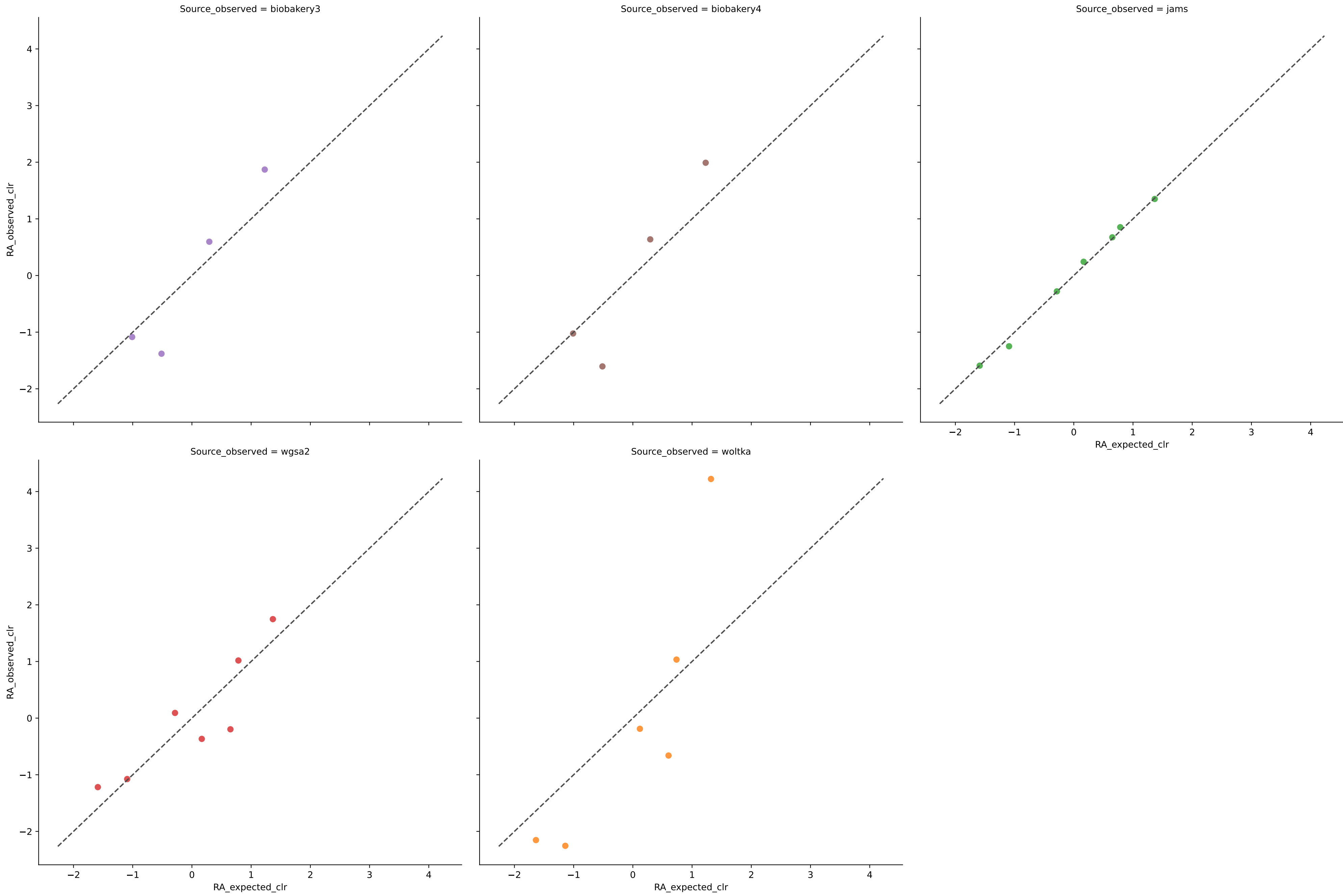


	Diversity	R ²	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487937	18	0.0234	0.0058	3.2948	0.7376	0.0085	100.0000	0.0000
SRR11487938	18	0.0204	0.0058	3.3182	0.7373	0.0085	100.0000	0.0000
SRR11487939	18	0.0221	0.0058	3.3334	0.7377	0.0084	100.0000	0.0000
SRR11487940	18	0.0196	0.0058	3.2964	0.7397	0.0084	100.0000	0.0000
SRR11487941	18	0.0193	0.0058	3.2935	0.7407	0.0084	100.0000	0.0000
Average	18	0.0210	0.0058	3.3073	0.7386	0.0085	100.0000	0.0000

Bivariate Linear Regression for Sample S1 in Experiment bmock12 (Genus at filter threshold 0.001)

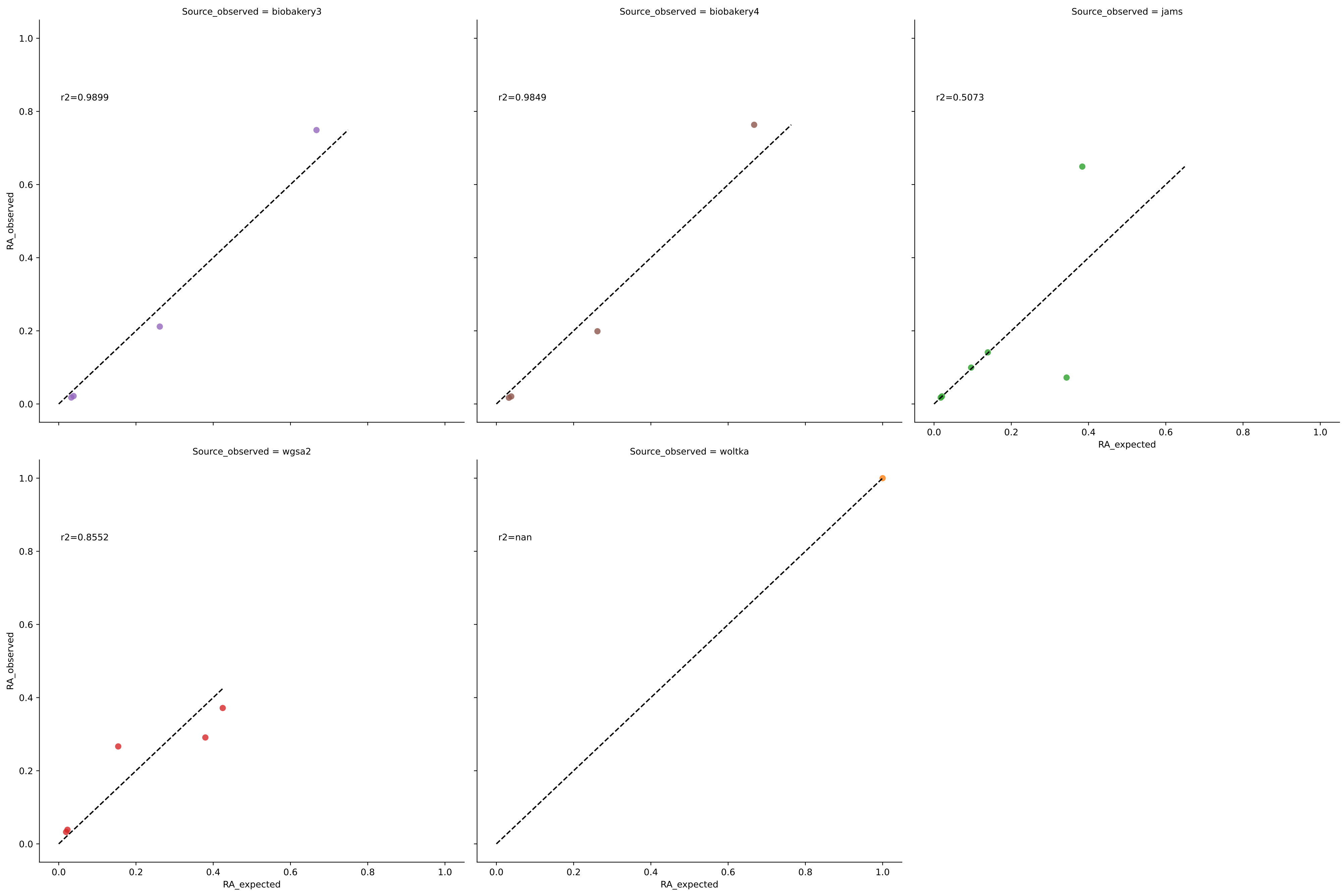


Bivariate Linear Regression for Sample S1 in Experiment bmock12 (Genus at filter threshold 0.001)

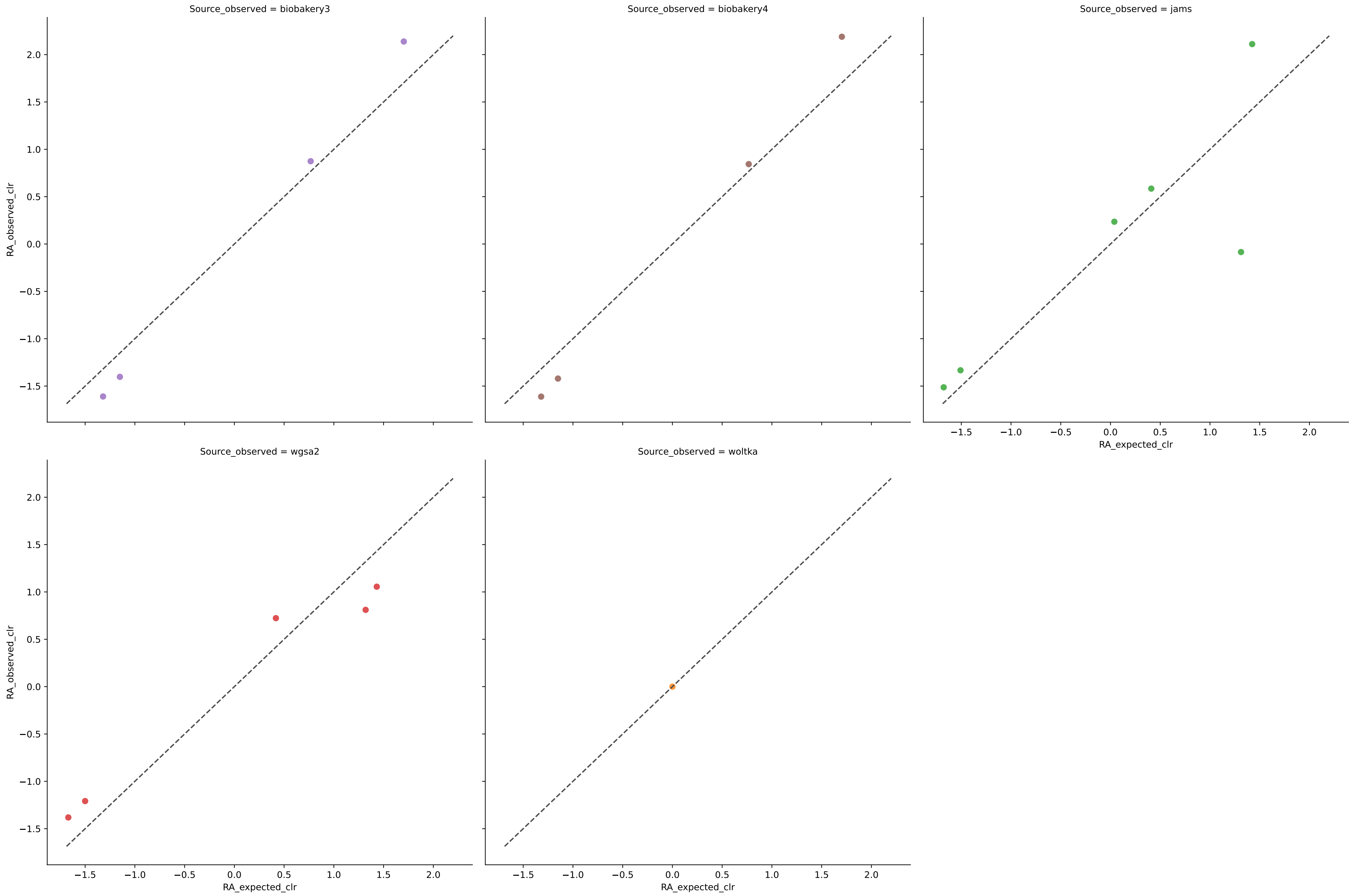


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
biobakery3	4	0.9911	0.0661	1.1219	0.8679	0.0788	100.0000	0.0000
biobakery4	4	0.9878	0.0757	1.3740	0.8486	0.0899	100.0000	0.0000
jams	7	0.9965	0.0053	0.1890	0.9813	0.0070	100.0000	0.0000
wgsa2	7	0.8585	0.0483	1.2171	0.8311	0.0654	100.0000	0.0000
woltka	6	0.6881	0.1791	3.4215	0.4627	0.2492	100.0000	0.0000

Bivariate Linear Regression for Sample S1 in Experiment bmock12 (Species at filter threshold 0.001)

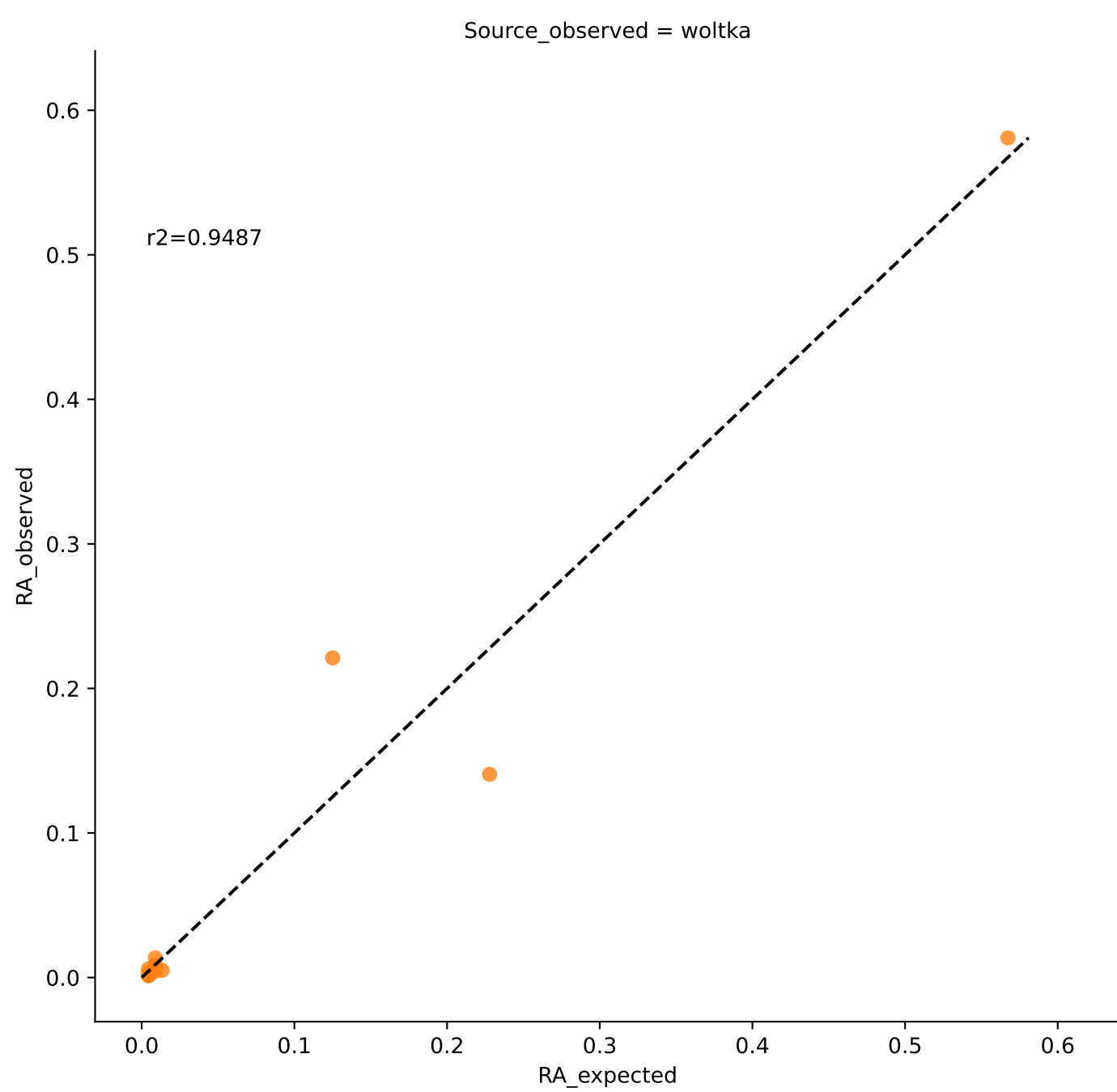
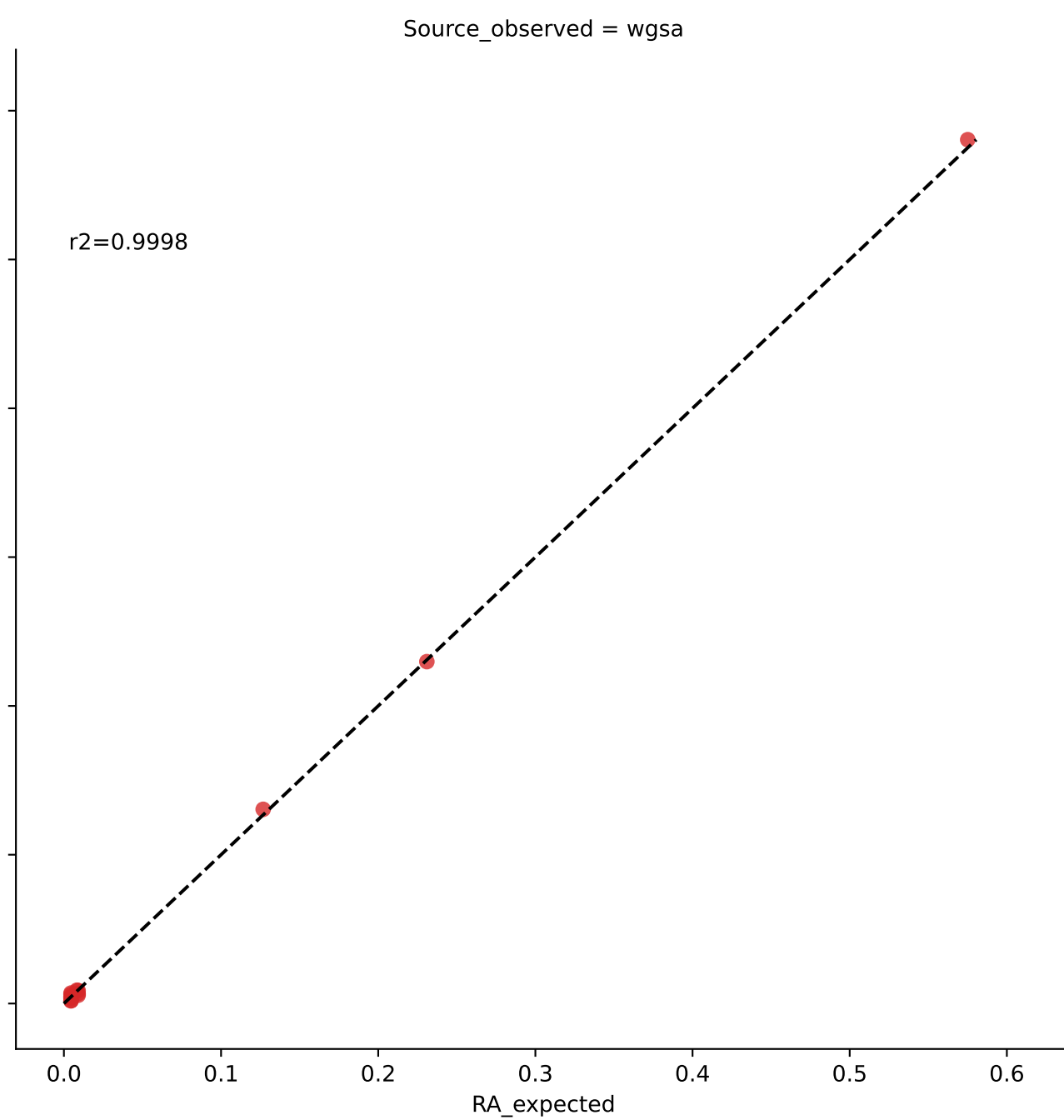
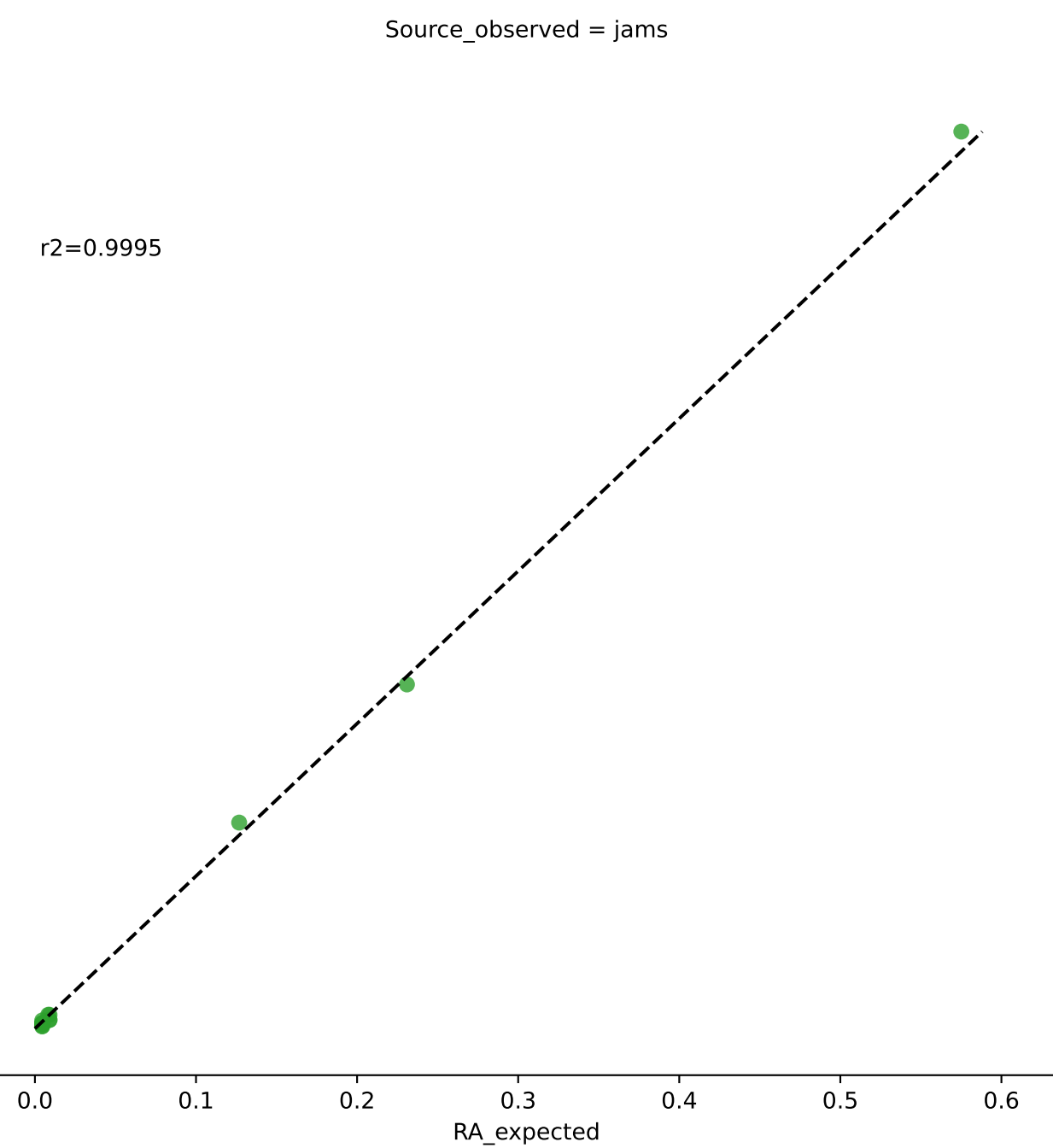
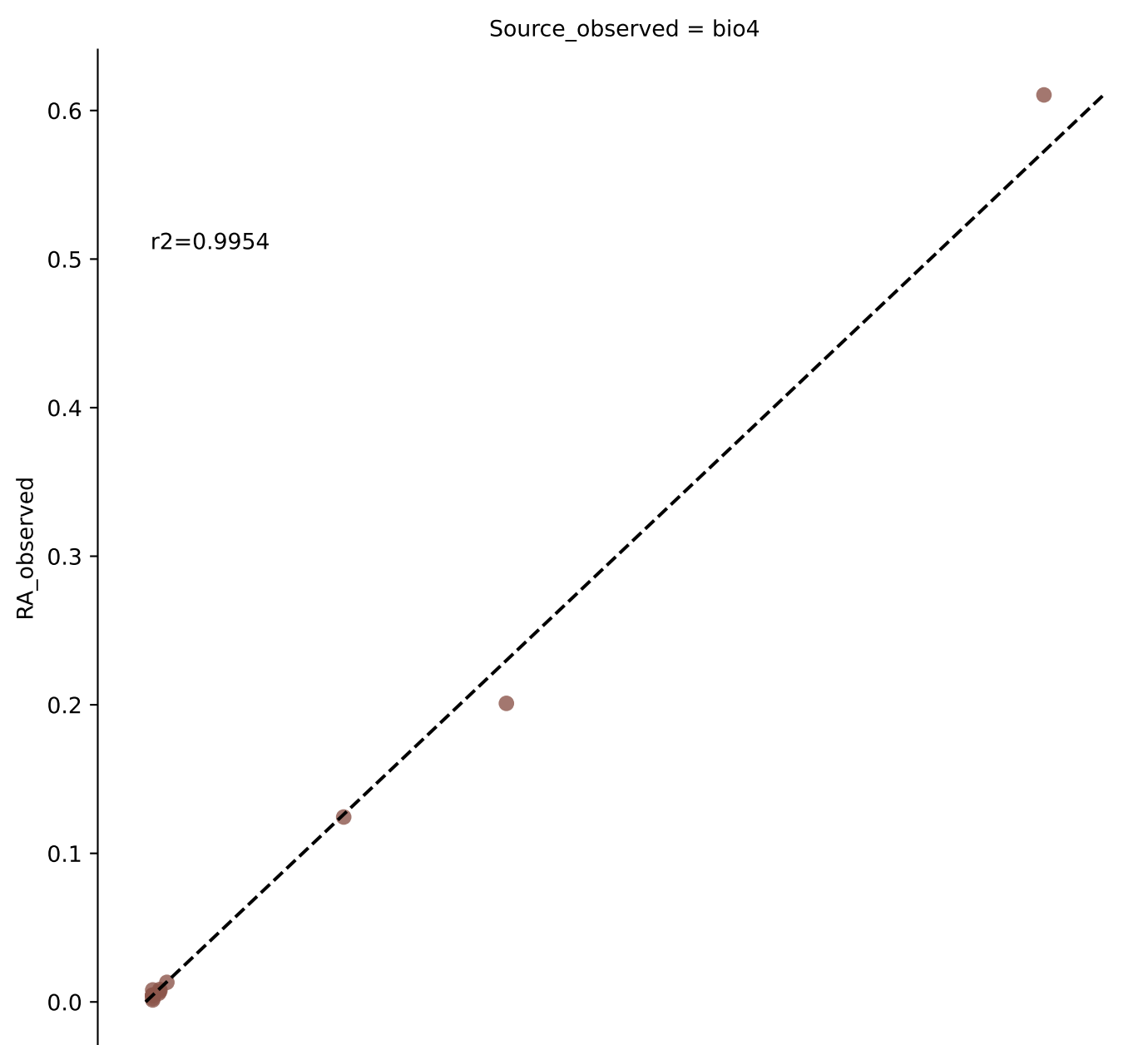


Bivariate Linear Regression for Sample S1 in Experiment bmock12 (Species at filter threshold 0.001)

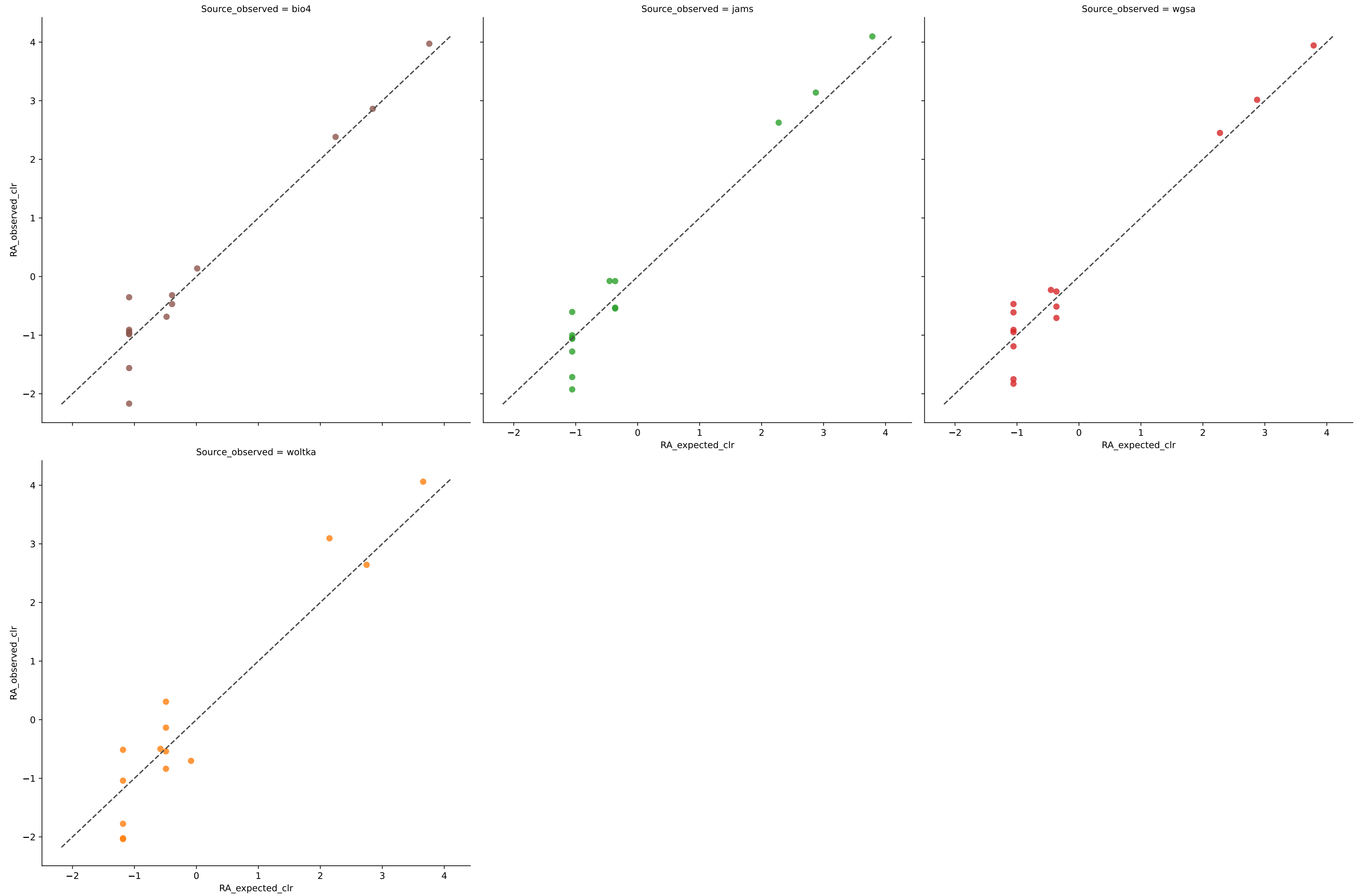


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
biobakery3	4	0.9899	0.0408	0.5907	0.9184	0.0492	100.0000	0.0000
biobakery4	4	0.9849	0.0480	0.6324	0.9040	0.0586	100.0000	0.0000
jams	6	0.5073	0.0903	1.5970	0.7290	0.1548	100.0000	0.0000
wgsa2	5	0.8552	0.0567	0.8135	0.8582	0.0690	100.0000	0.0000
woltka	1	nan	0.0000	0.0000	1.0000	0.0000	100.0000	0.0000

Bivariate Linear Regression for Sample S1 in Experiment camisimGI (Genus at filter threshold 0.001)

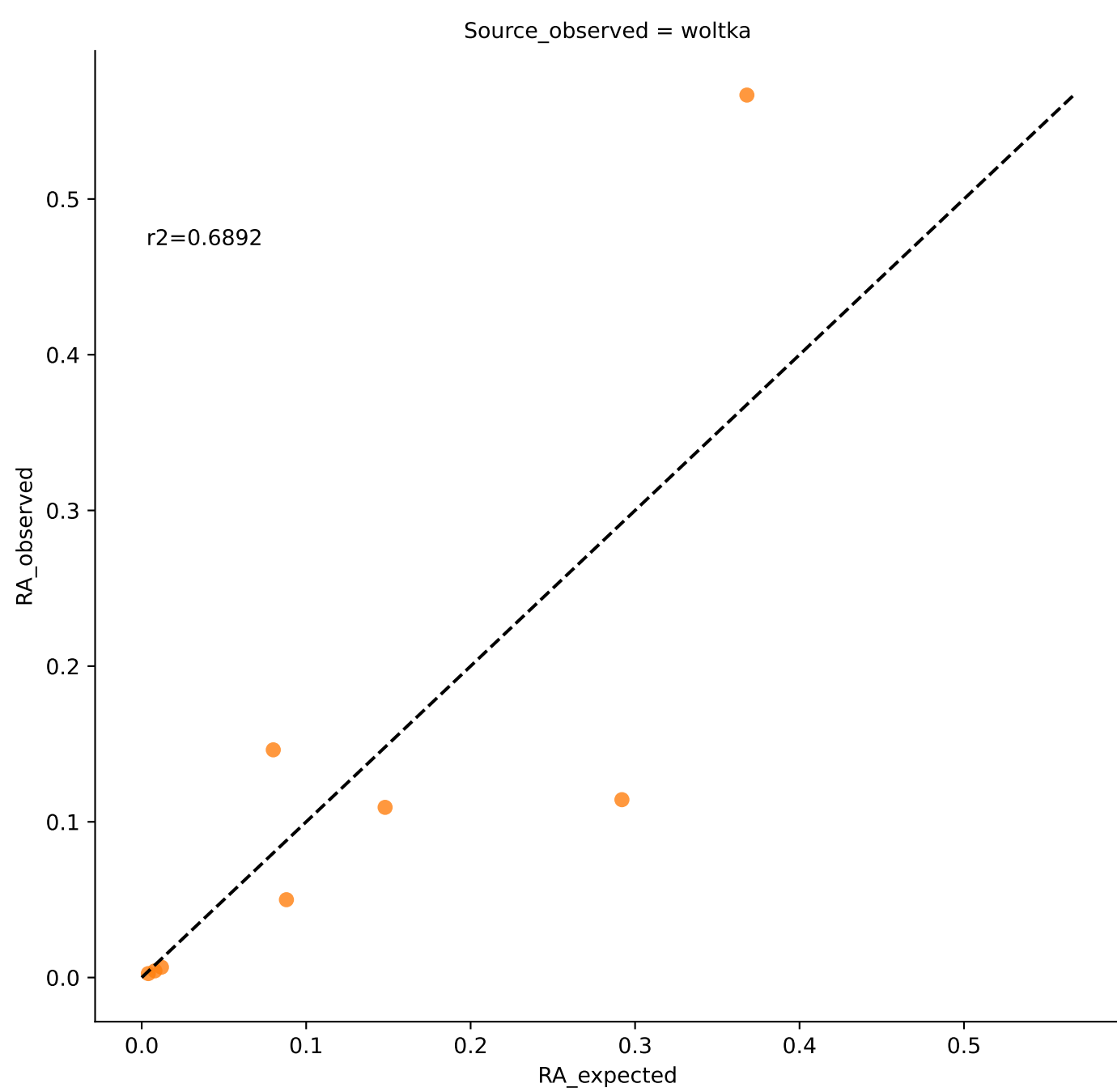
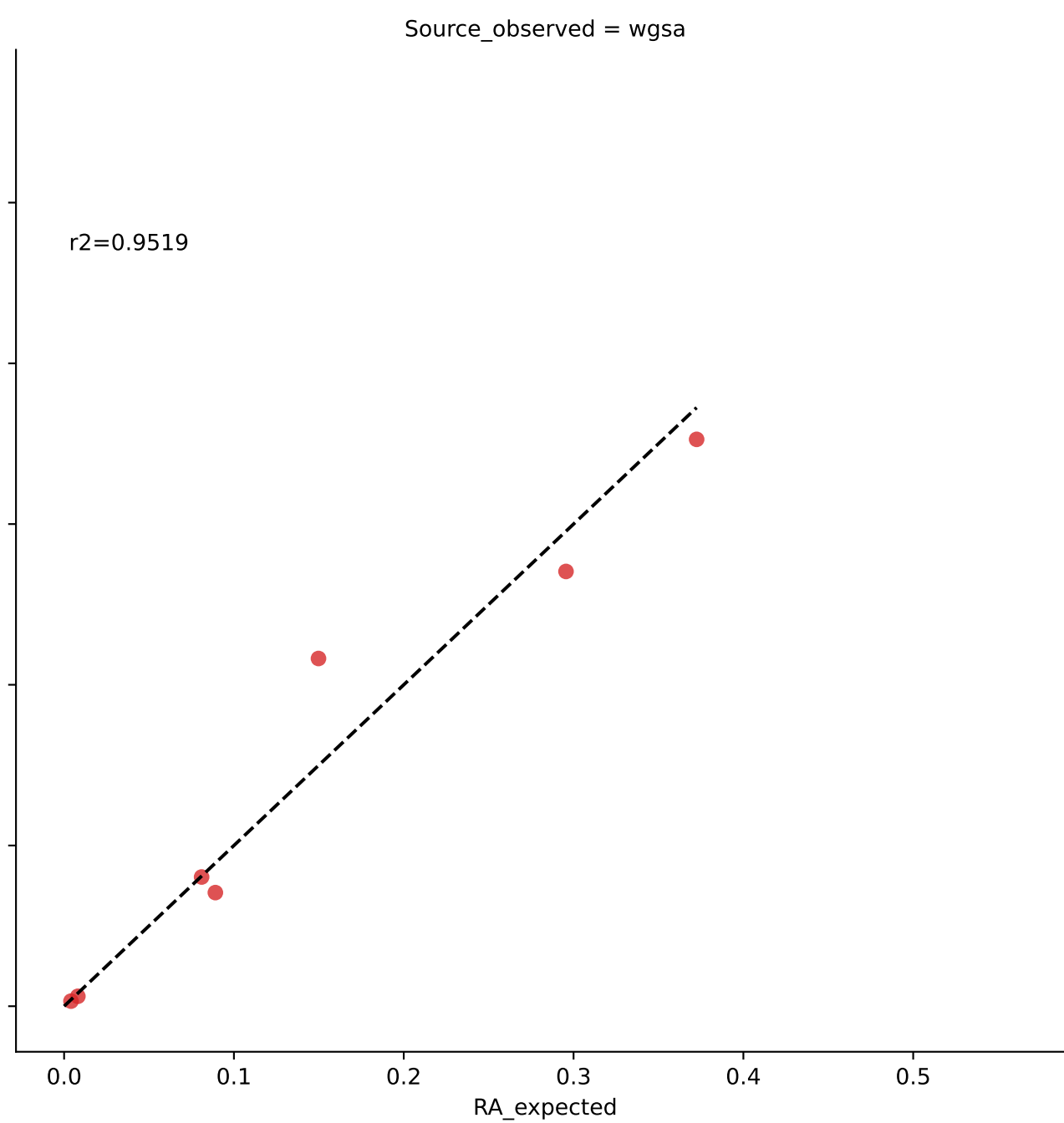
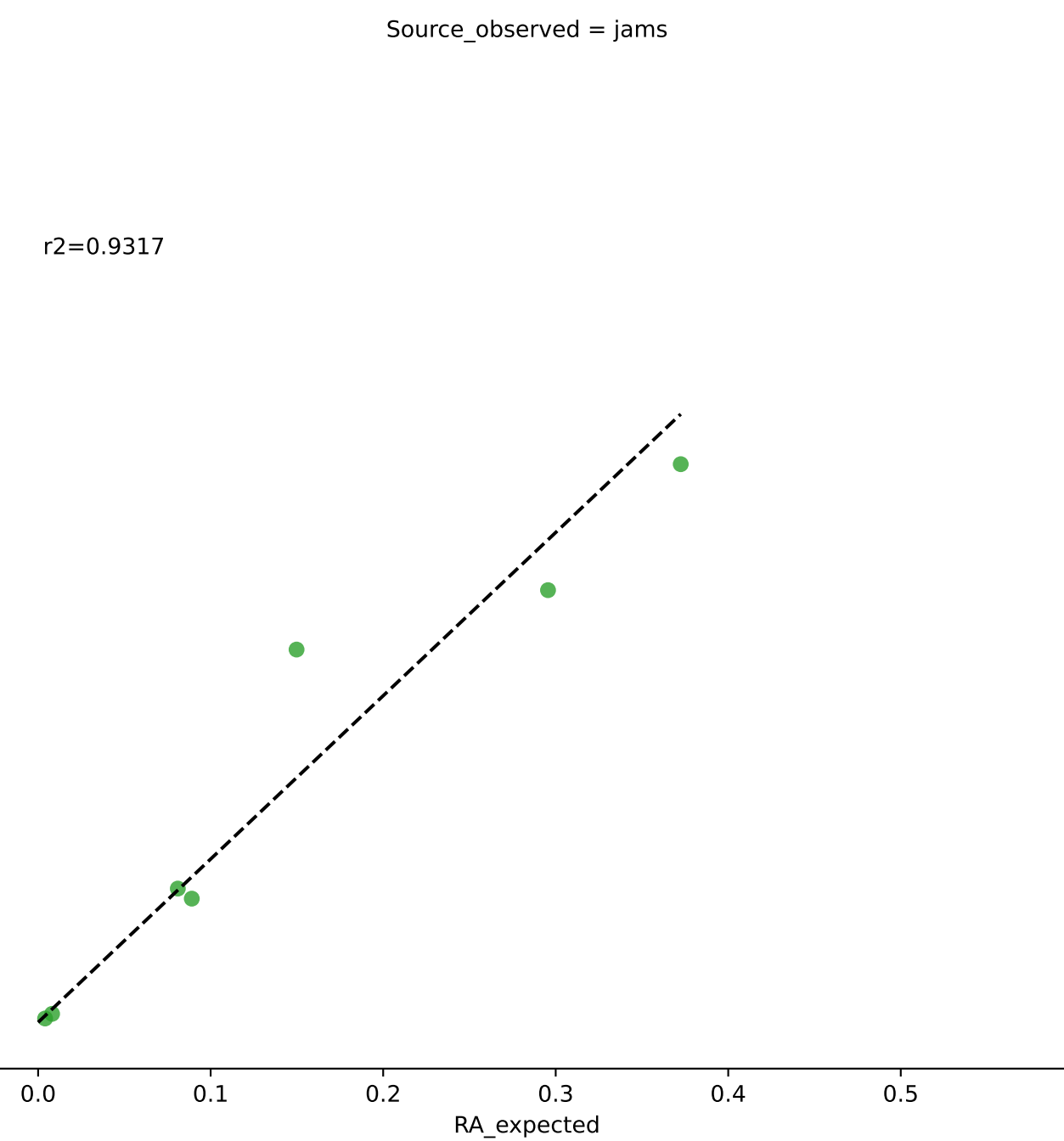
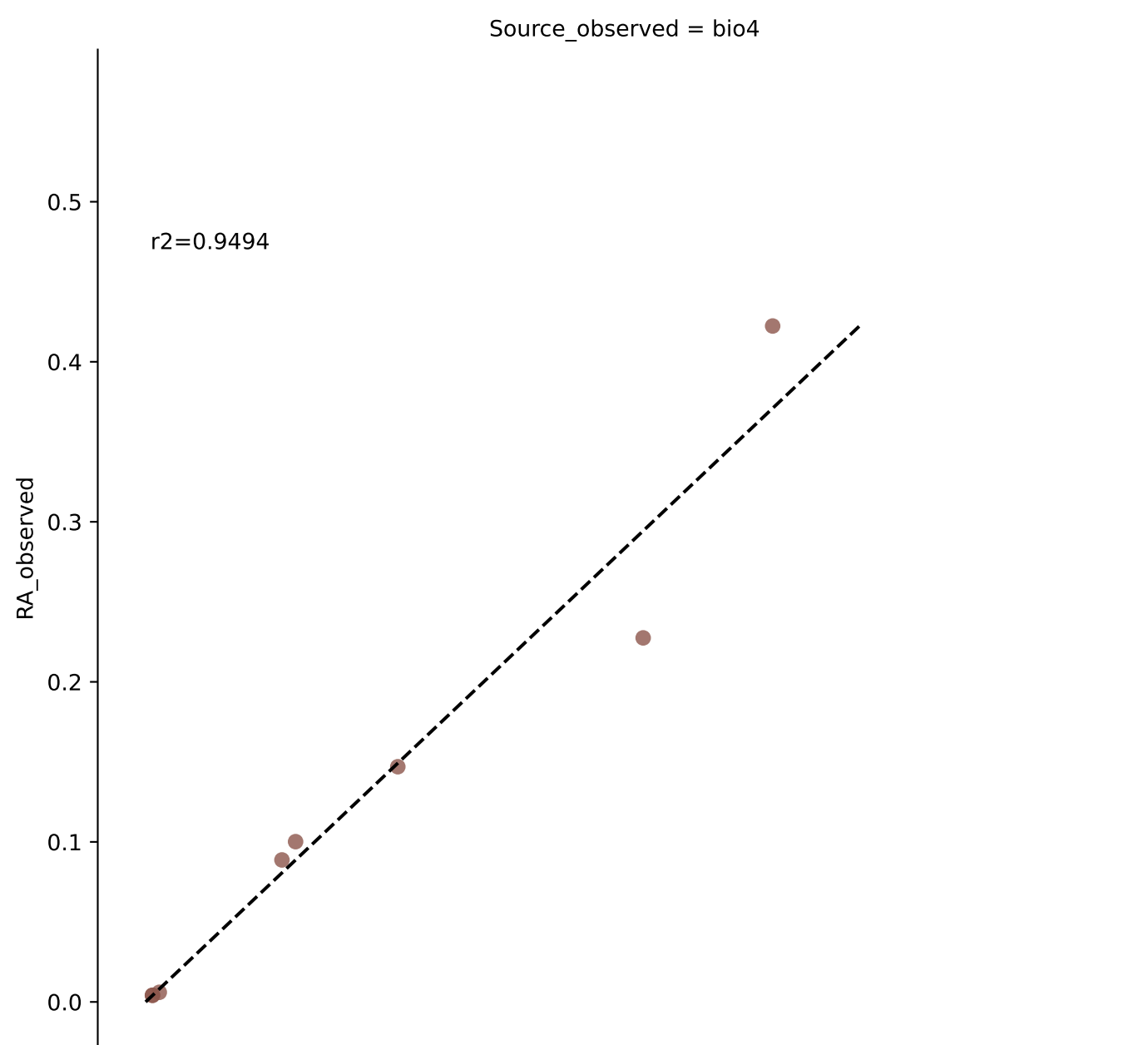


Bivariate Linear Regression for Sample S1 in Experiment camisimGI (Genus at filter threshold 0.001)

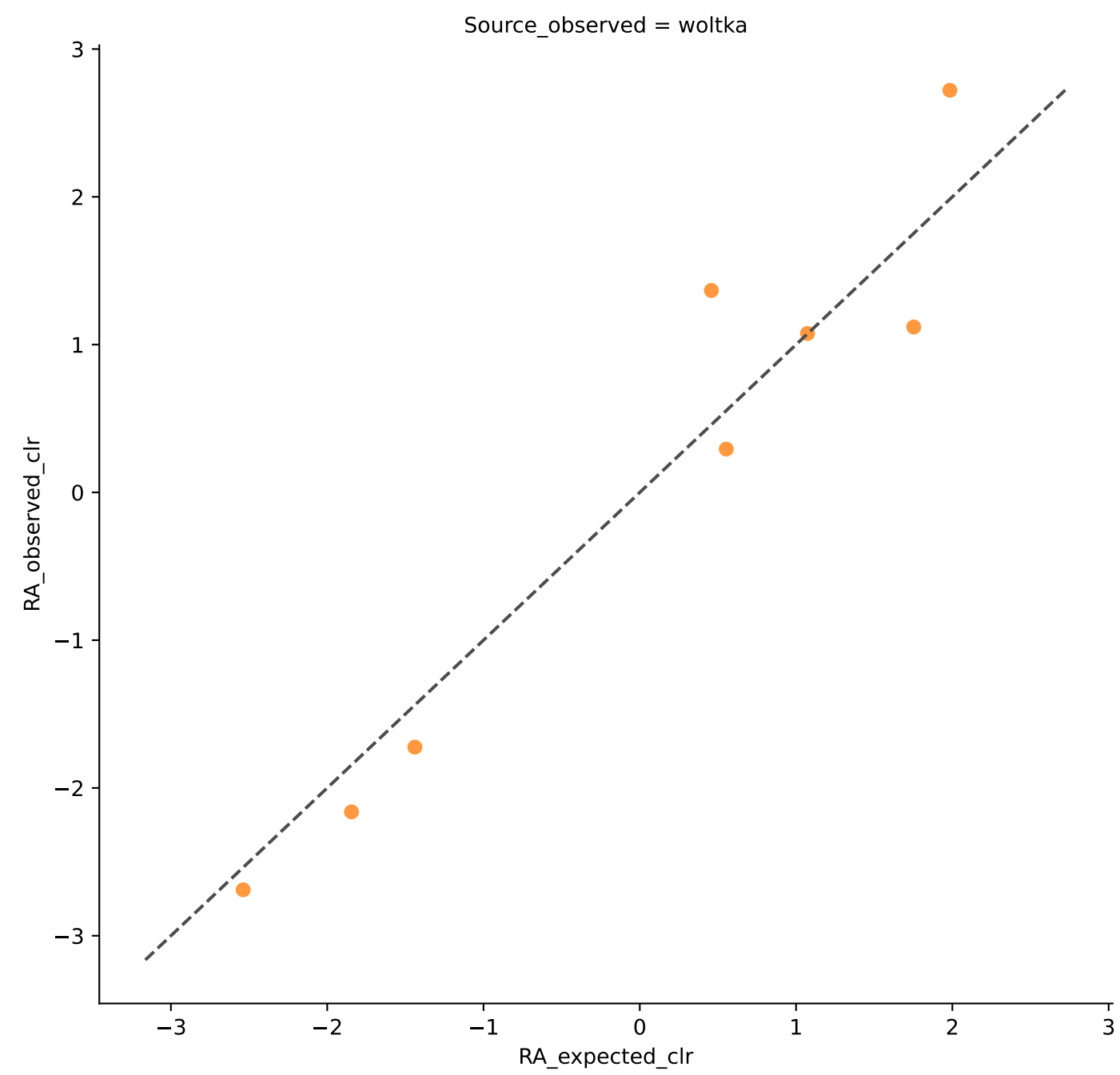
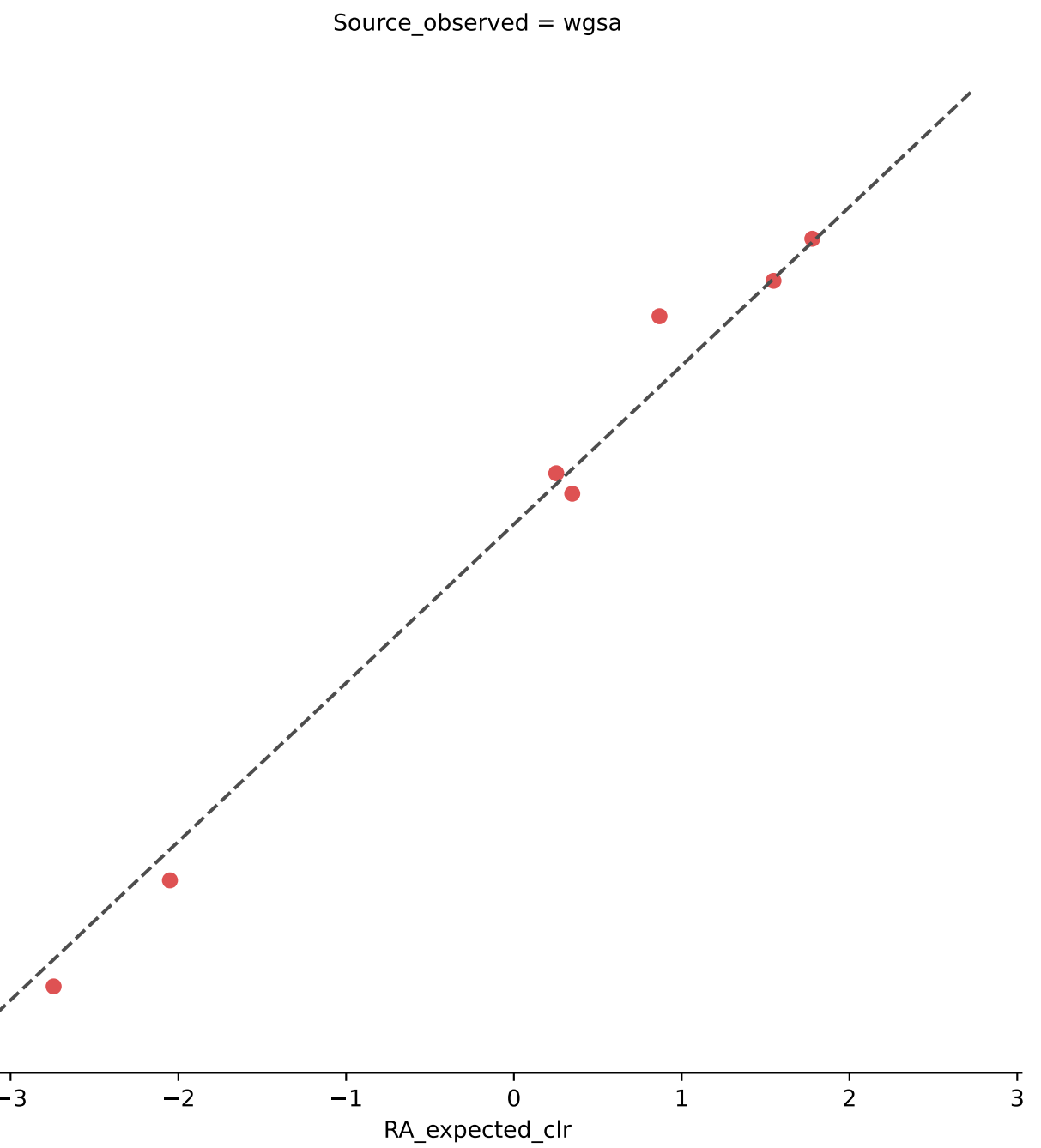
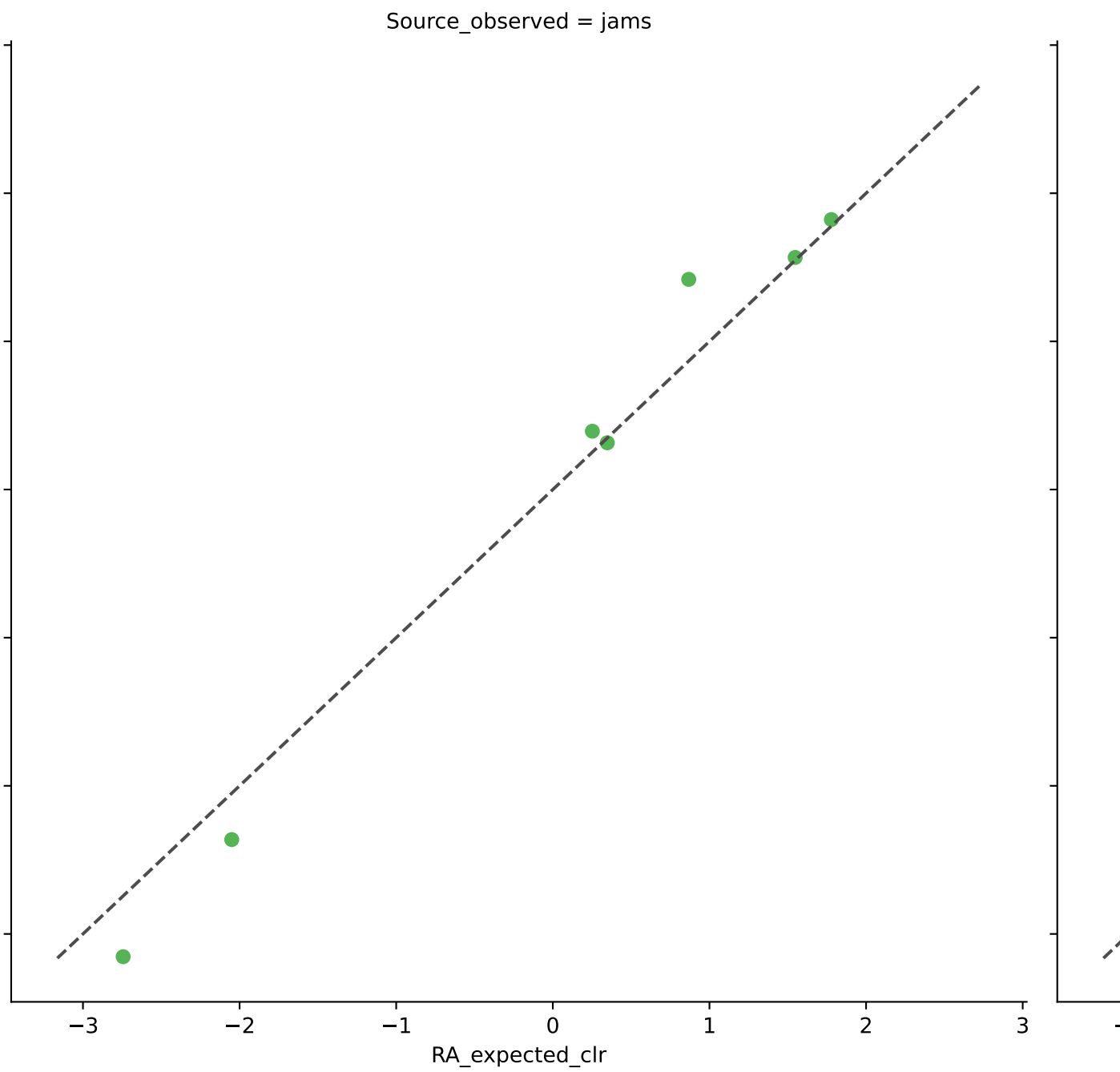
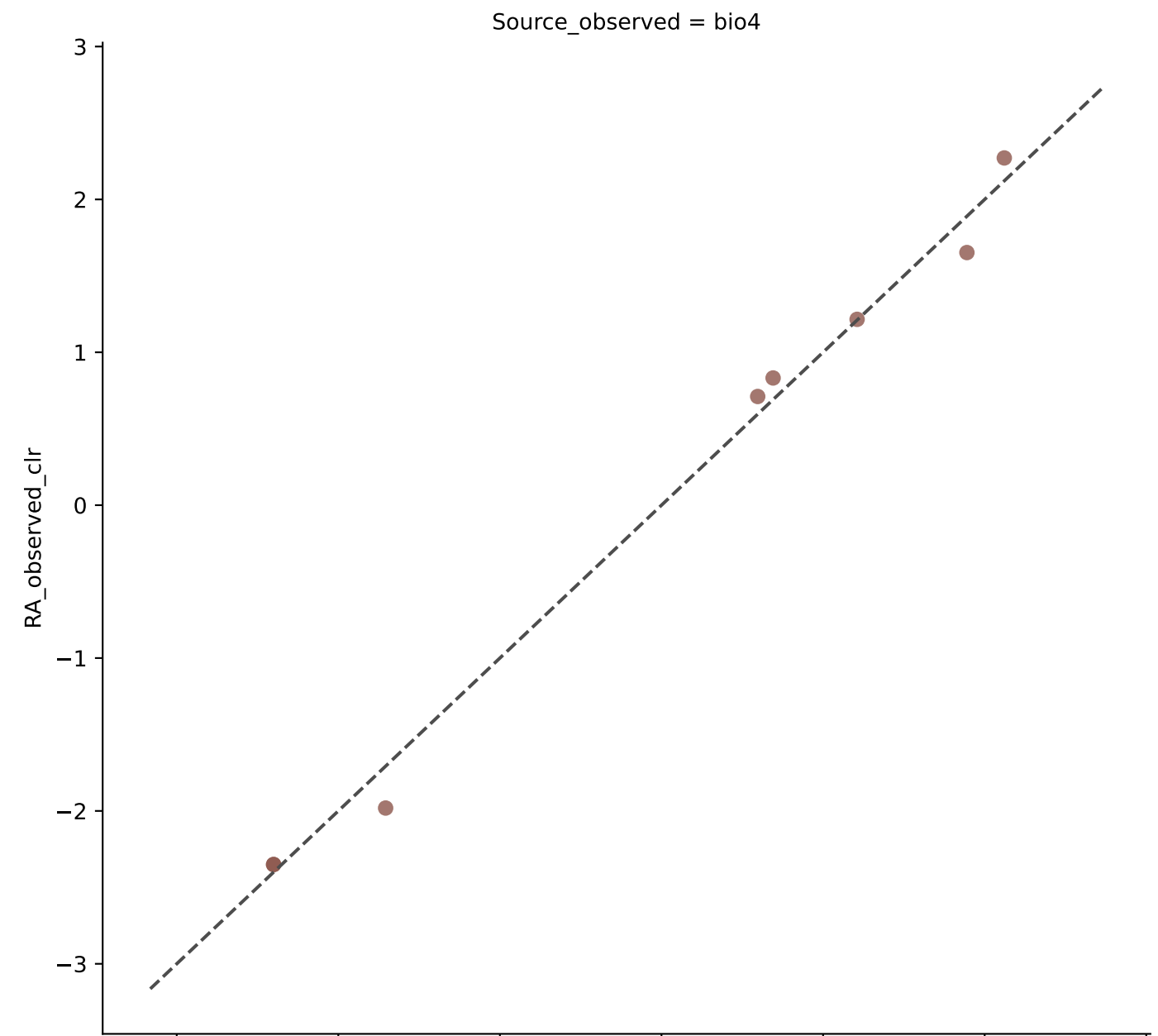


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	14	0.9954	0.0060	1.4642	0.9583	0.0129	100.0000	0.0000
jams	14	0.9995	0.0033	1.4210	0.9771	0.0047	100.0000	0.0000
wgsa	14	0.9998	0.0020	1.3962	0.9861	0.0025	100.0000	0.0000
woltka	14	0.9487	0.0165	2.1427	0.8843	0.0350	100.0000	0.0000

Bivariate Linear Regression for Sample S2 in Experiment camisimGI (Genus at filter threshold 0.001)

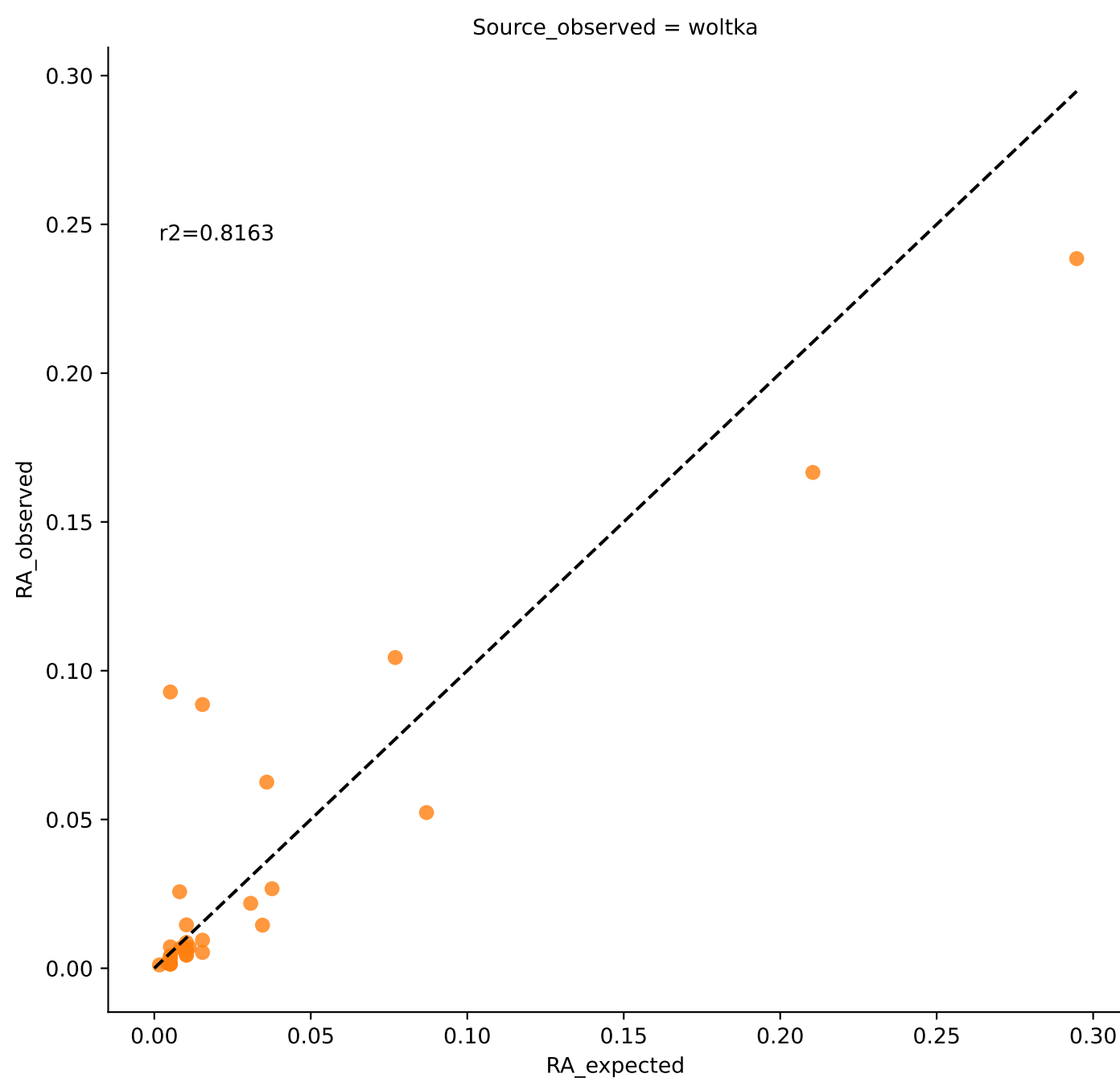
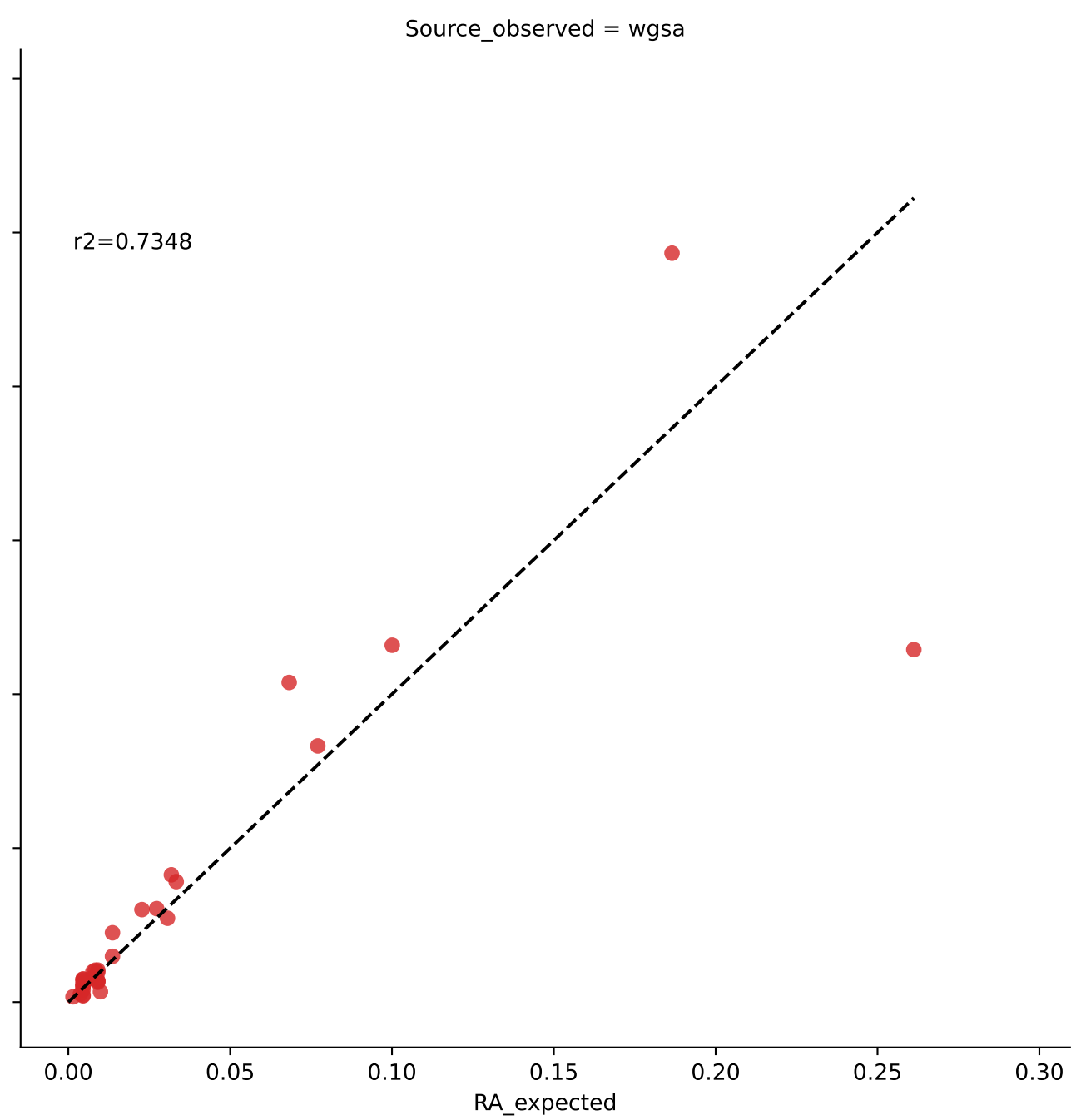
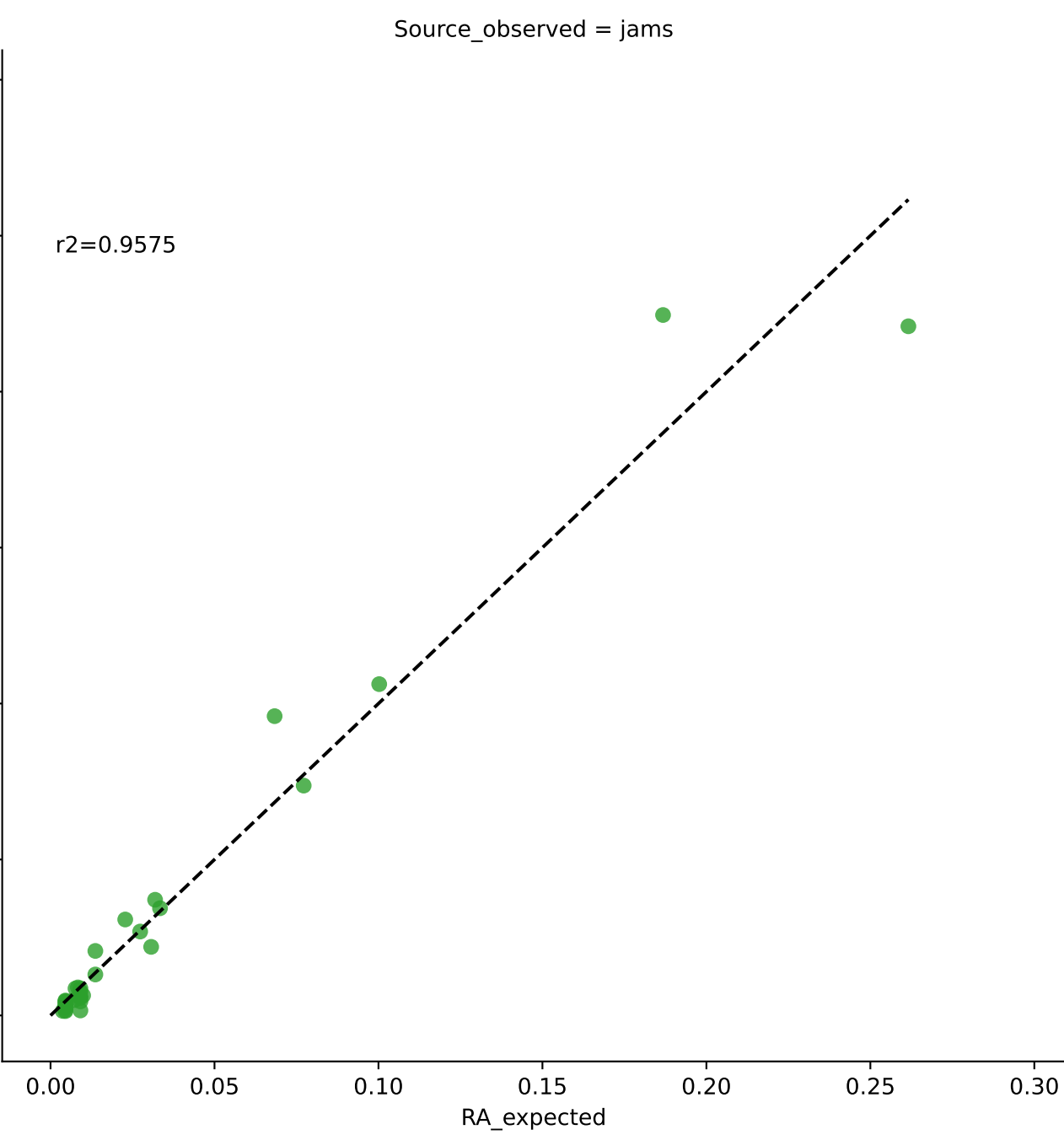
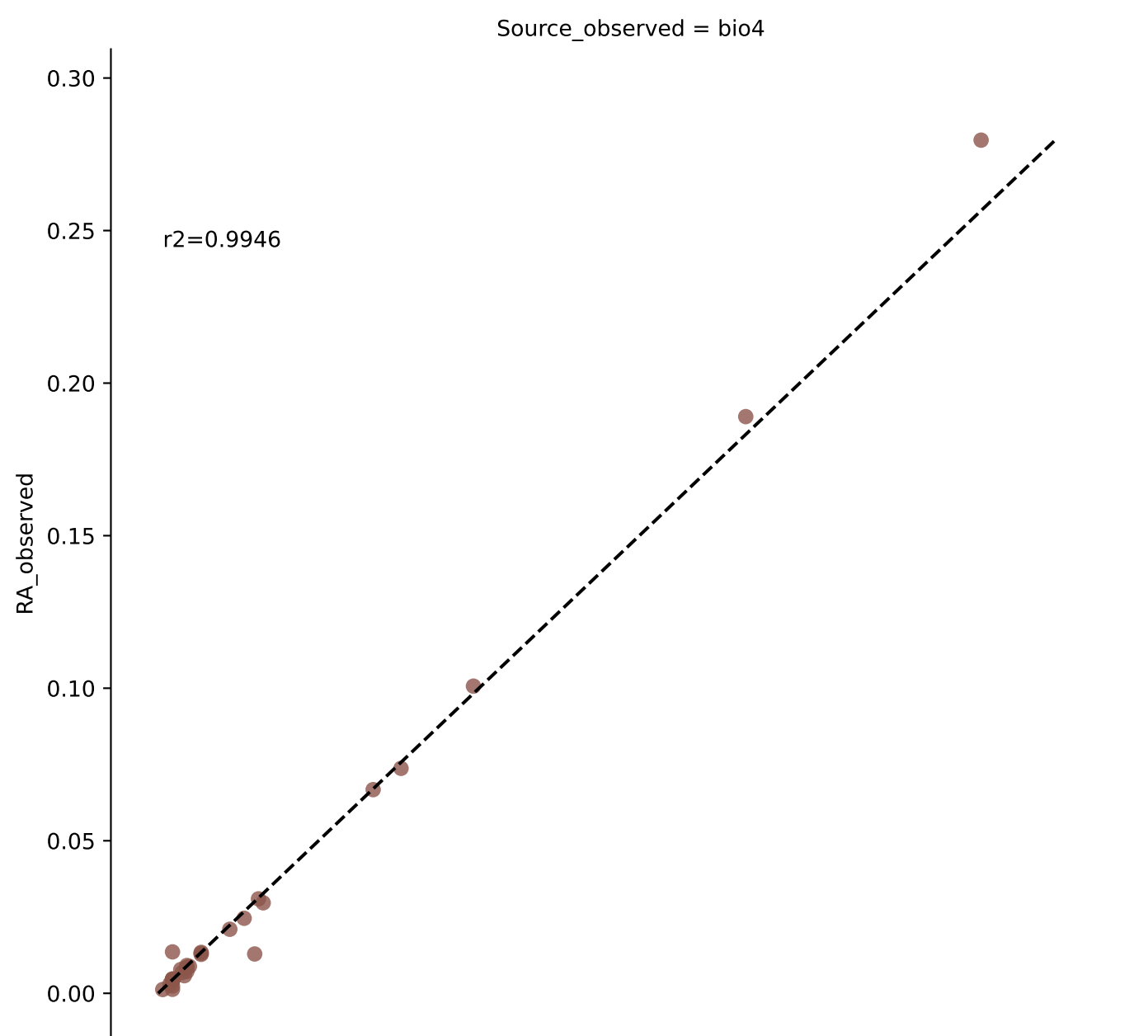


Bivariate Linear Regression for Sample S2 in Experiment camisimGI (Genus at filter threshold 0.001)

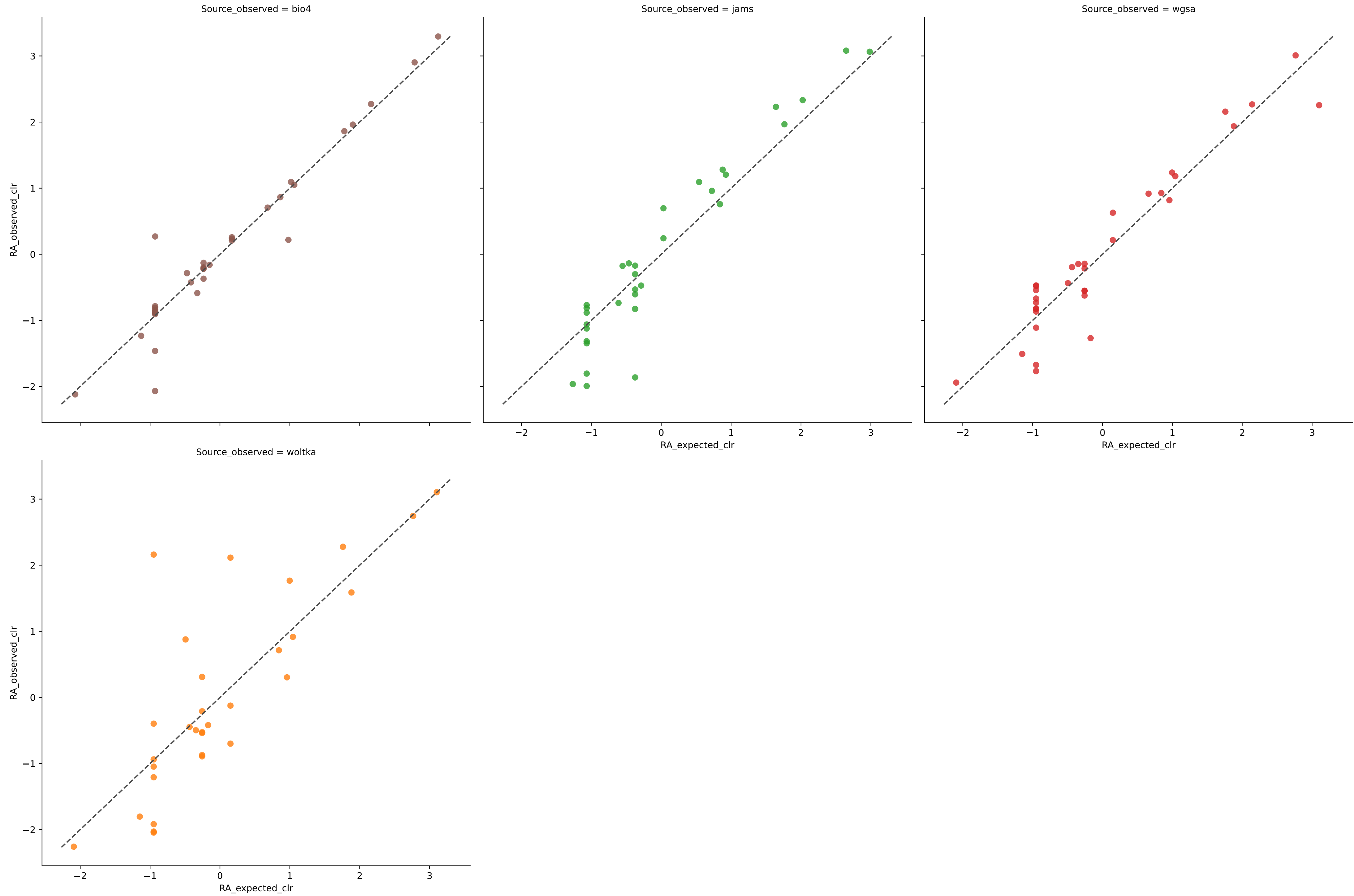


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	8	0.9494	0.0178	0.4385	0.9288	0.0302	100.0000	0.0000
jams	7	0.9317	0.0227	0.7701	0.9206	0.0343	100.0000	0.0000
wgsa	7	0.9519	0.0190	0.5411	0.9334	0.0288	100.0000	0.0000
woltka	8	0.6892	0.0662	1.4289	0.7350	0.0990	100.0000	0.0000

Bivariate Linear Regression for Sample S1 in Experiment camisimGI (Species at filter threshold 0.001)

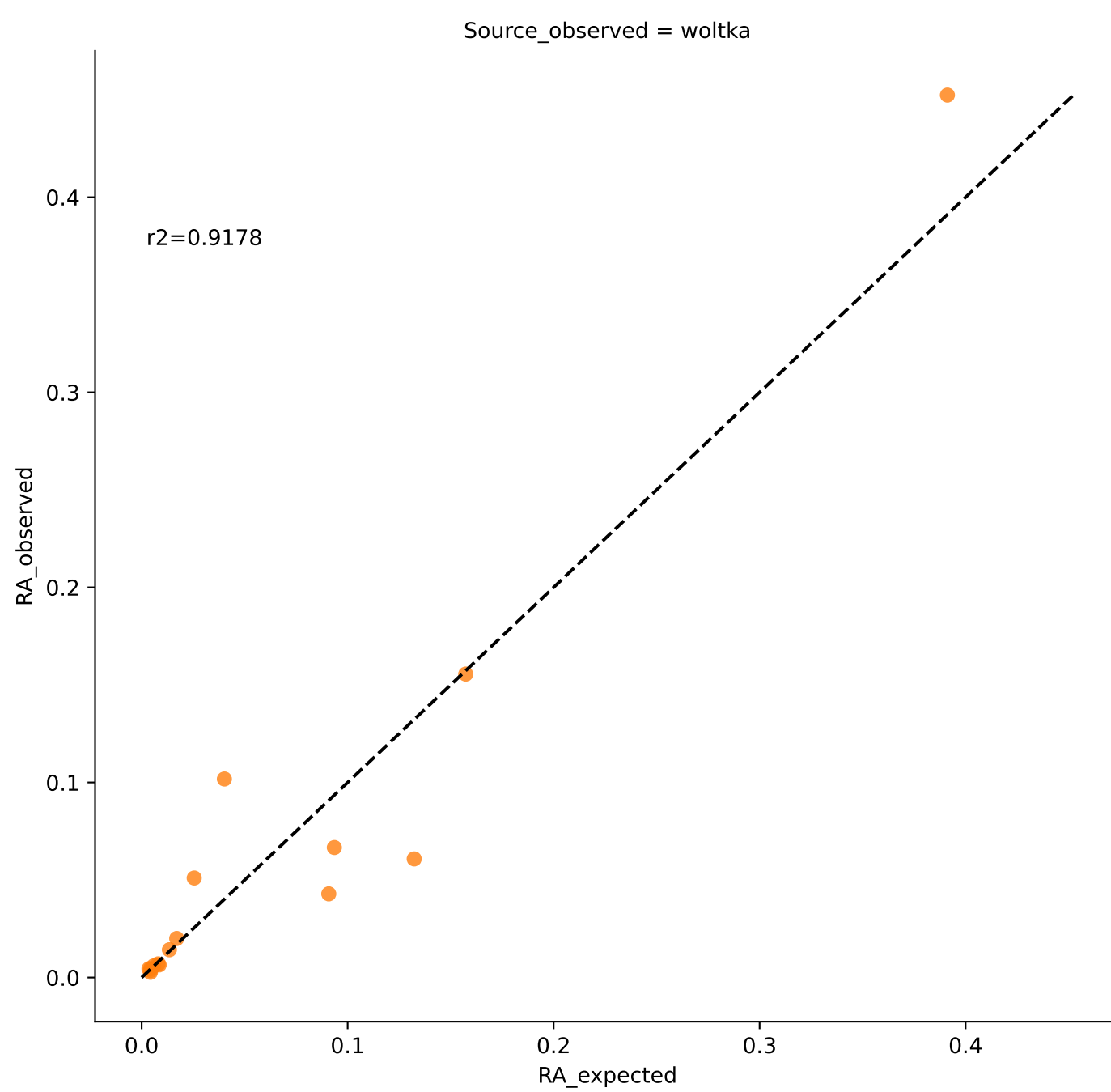
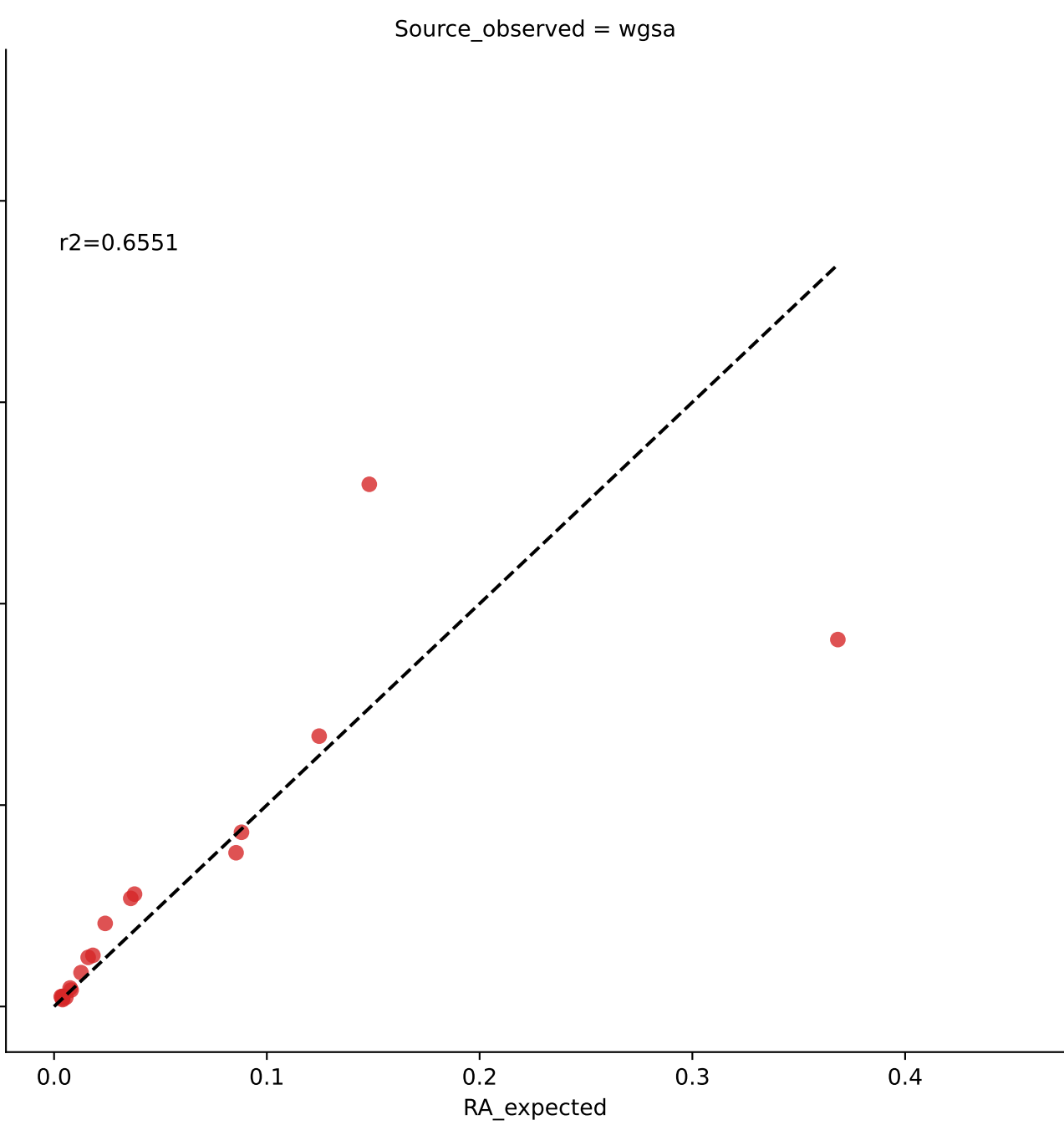
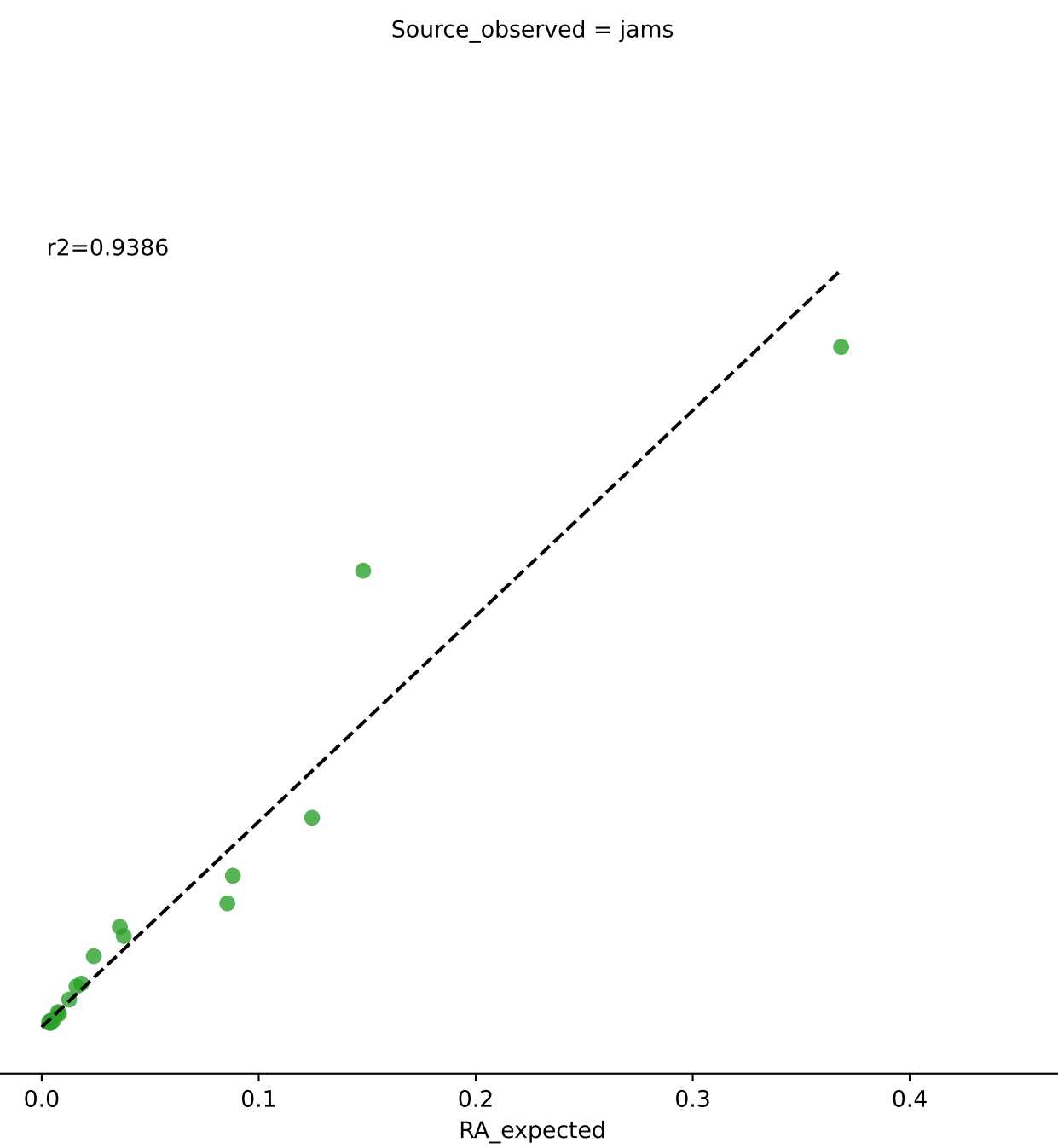
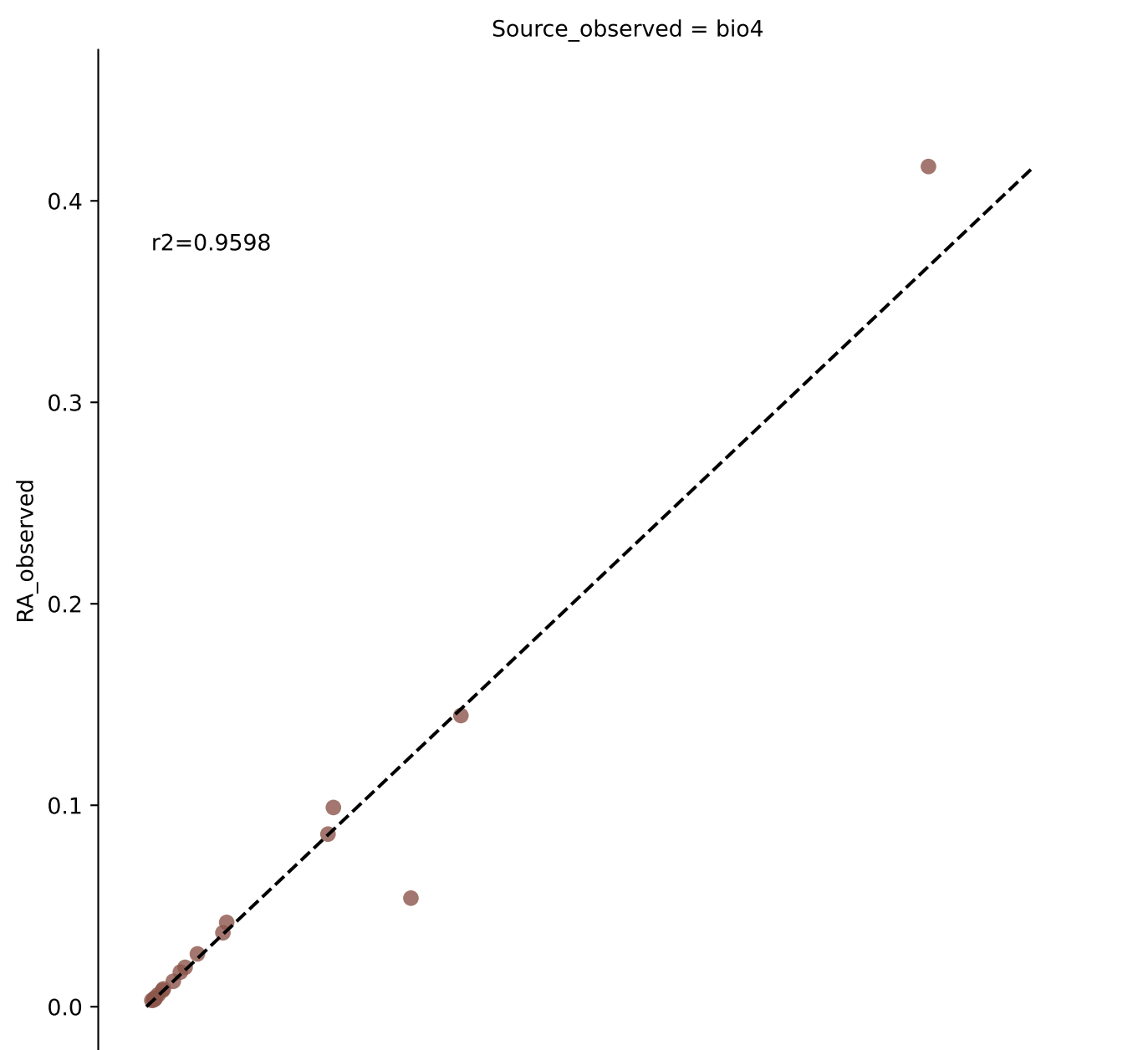


Bivariate Linear Regression for Sample S1 in Experiment camisimGI (Species at filter threshold 0.001)

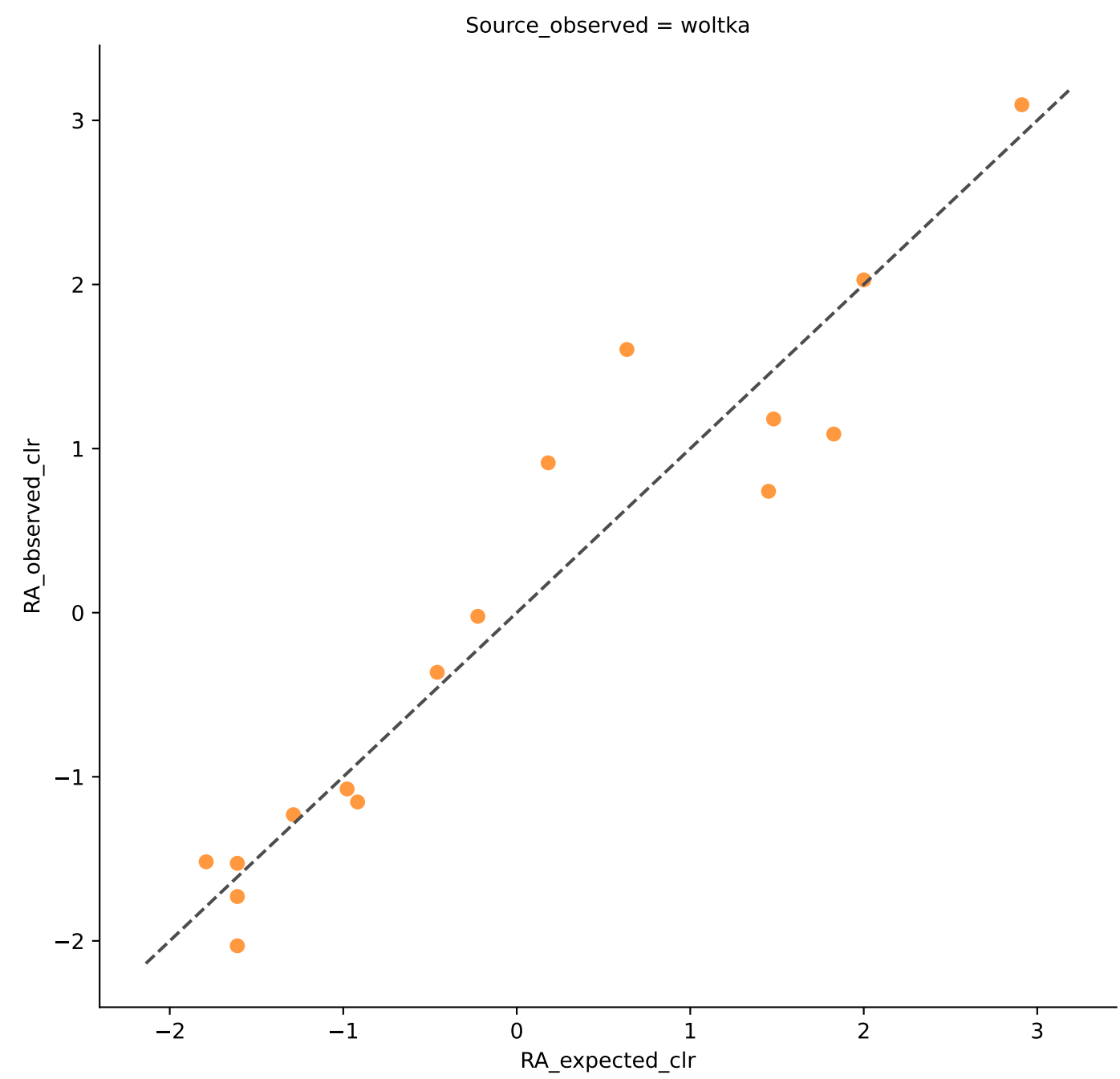
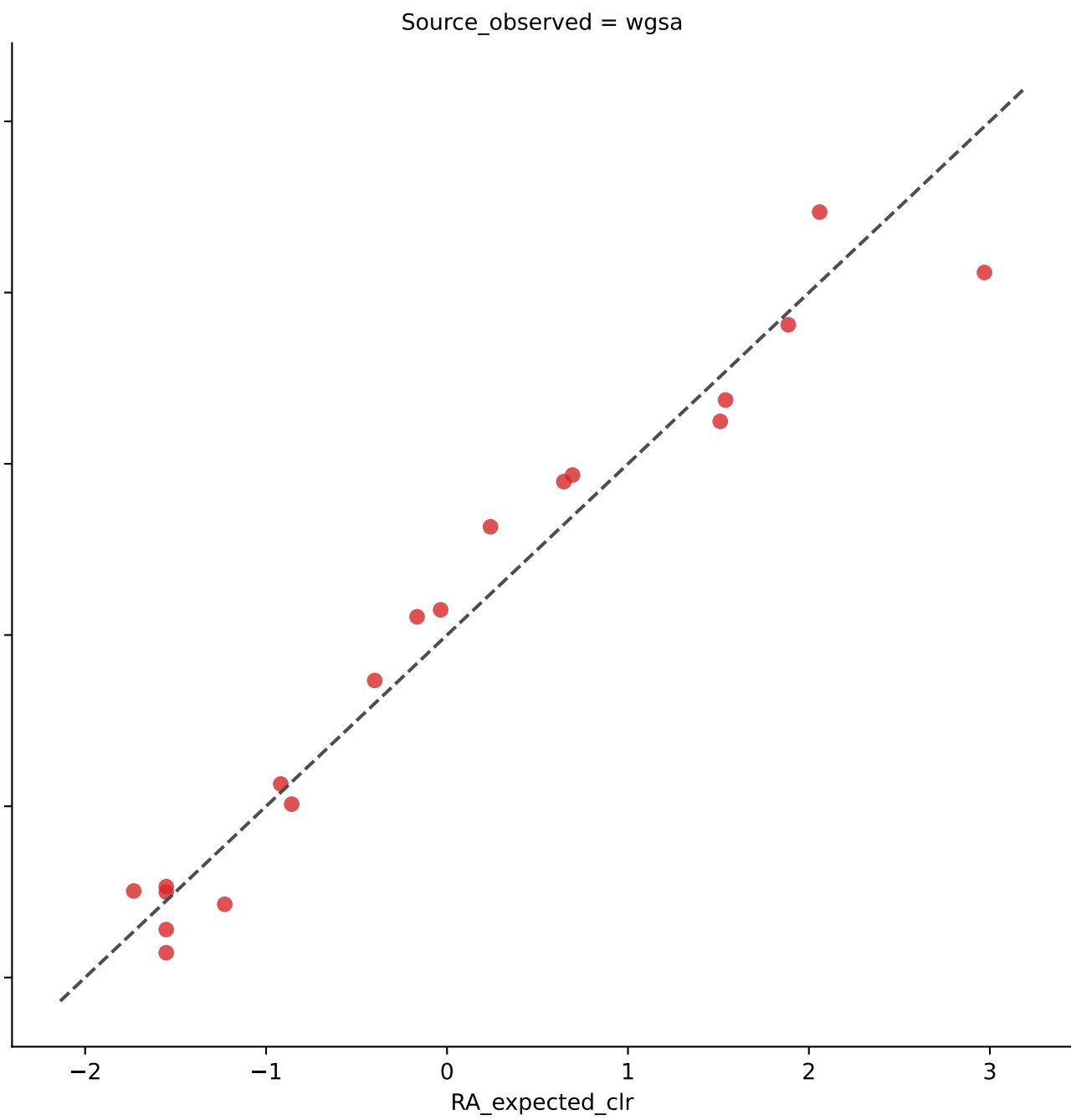
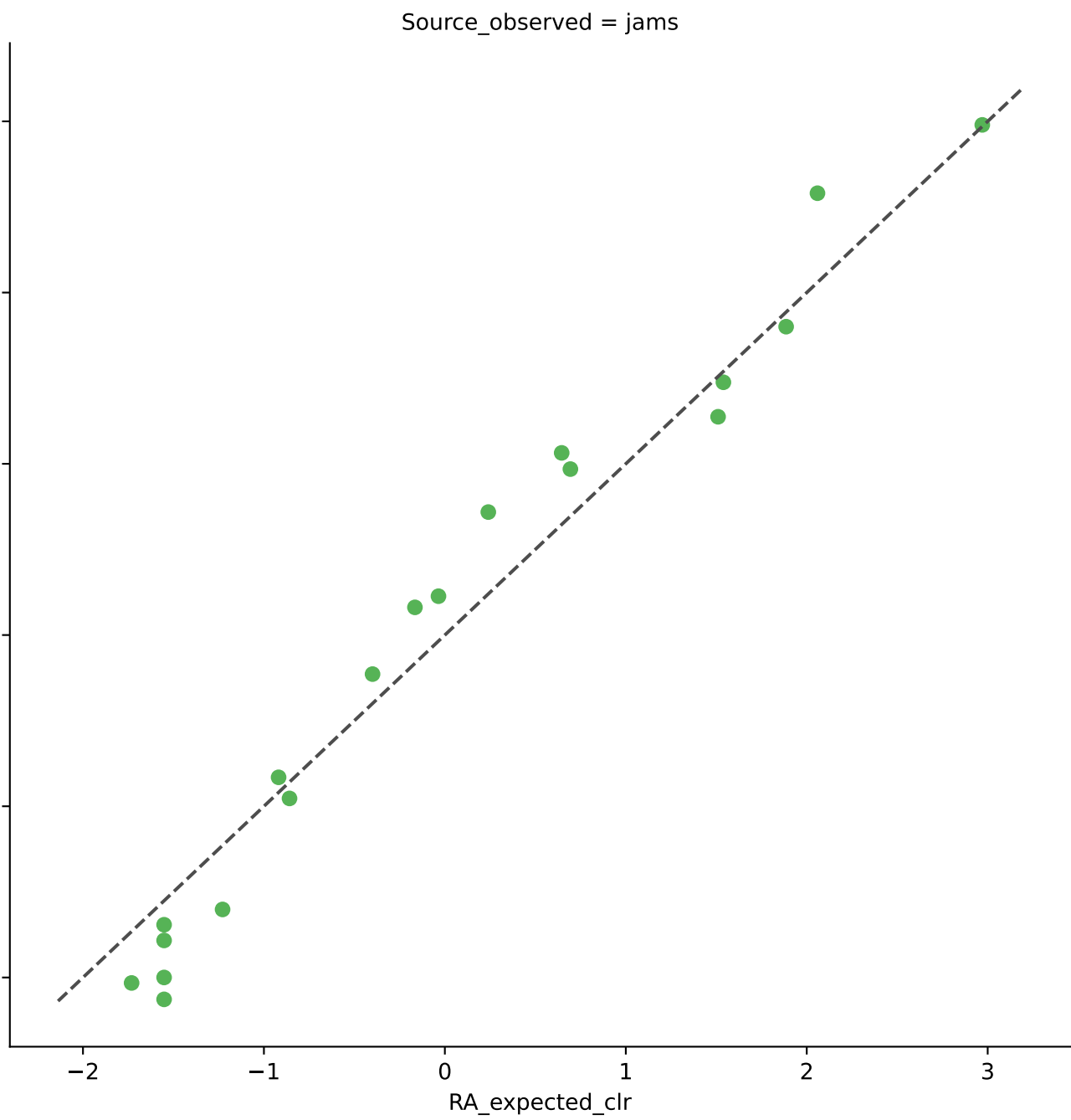
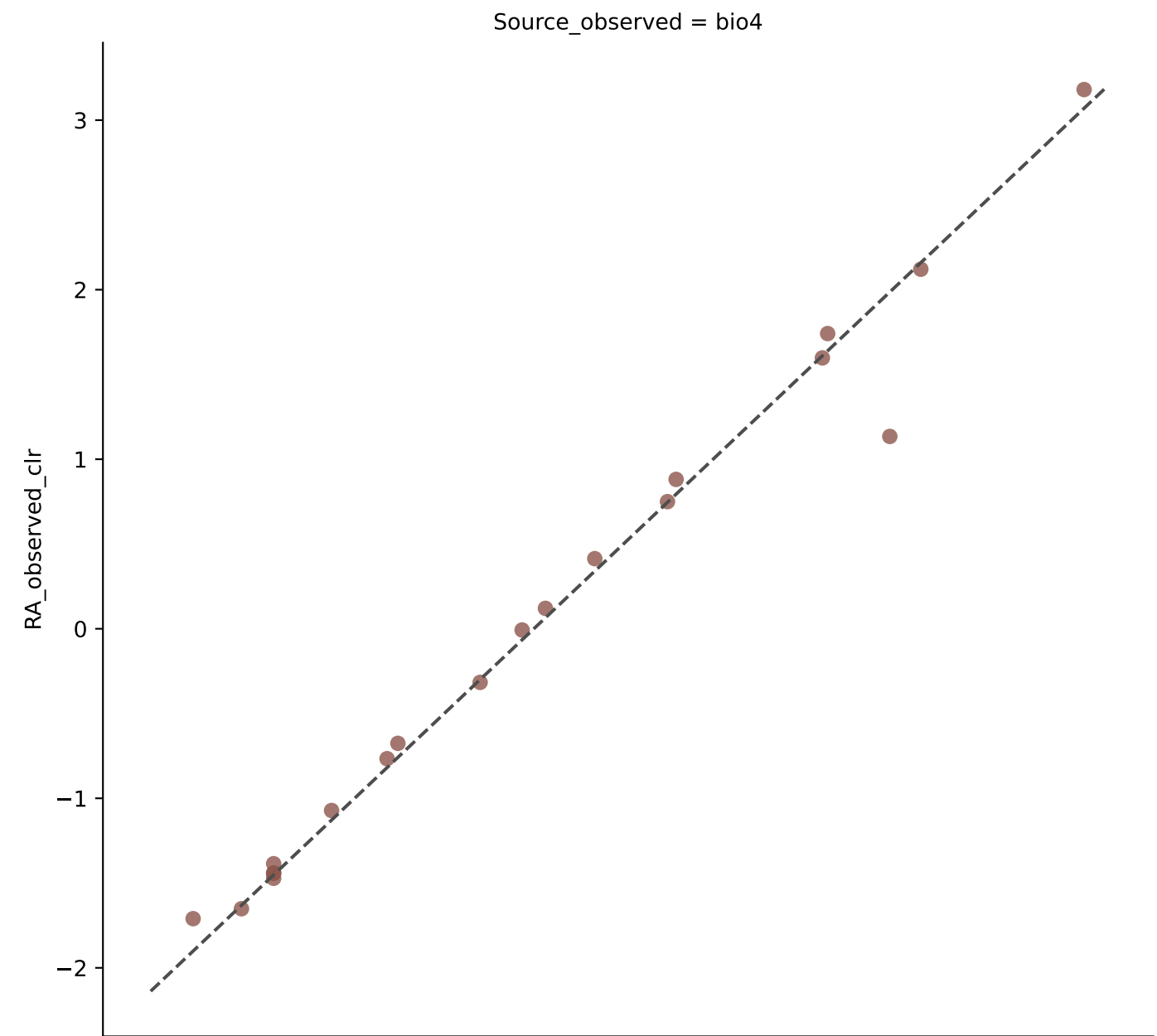


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	36	0.9946	0.0023	1.9734	0.9582	0.0053	100.0000	0.0000
jams	32	0.9575	0.0059	2.6148	0.9054	0.0115	100.0000	0.0000
wgsa	34	0.7348	0.0100	2.2676	0.8298	0.0280	100.0000	0.0000
woltka	31	0.8163	0.0154	4.8078	0.7610	0.0267	100.0000	0.0000

Bivariate Linear Regression for Sample S2 in Experiment camisimGI (Species at filter threshold 0.001)

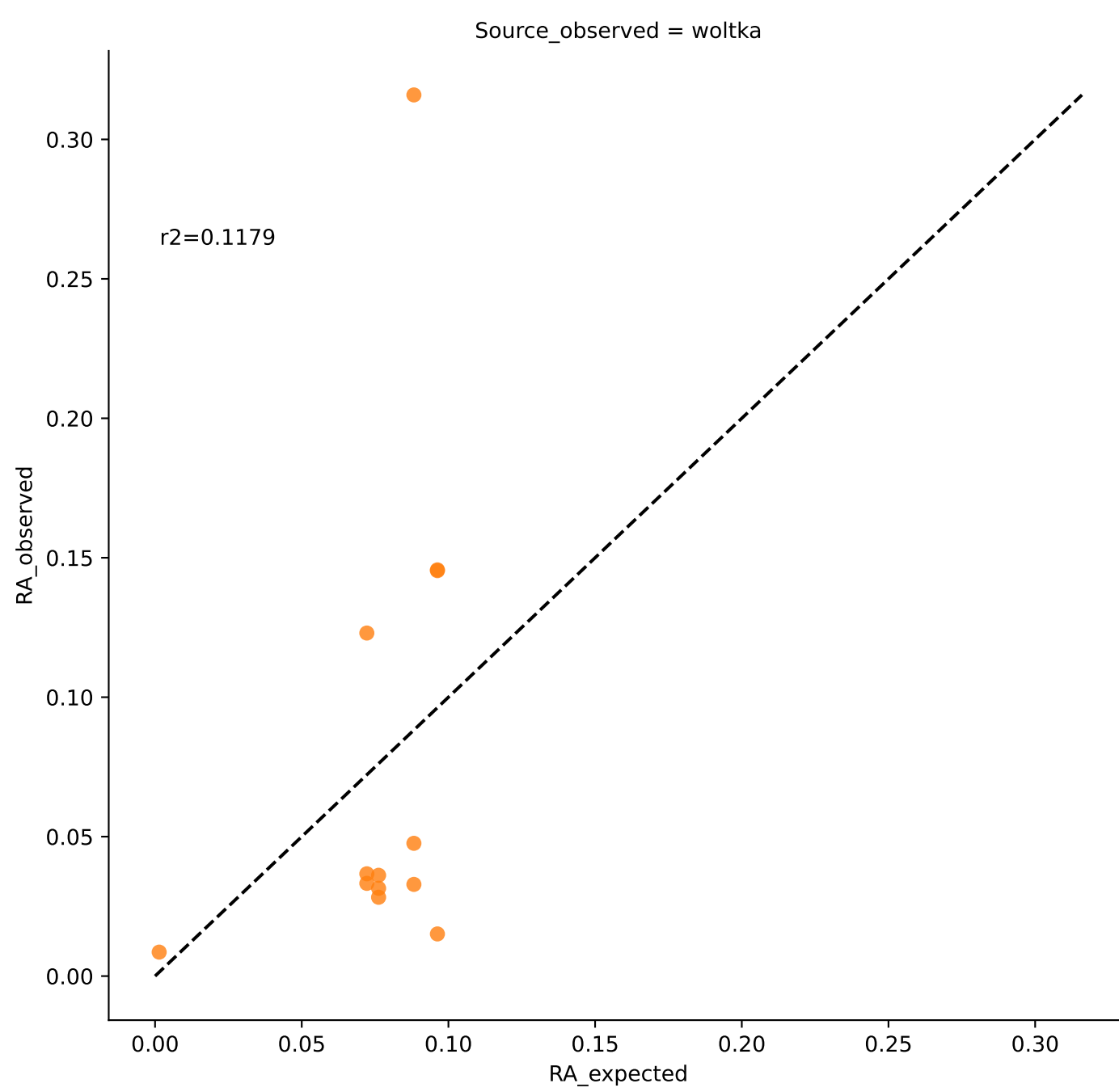
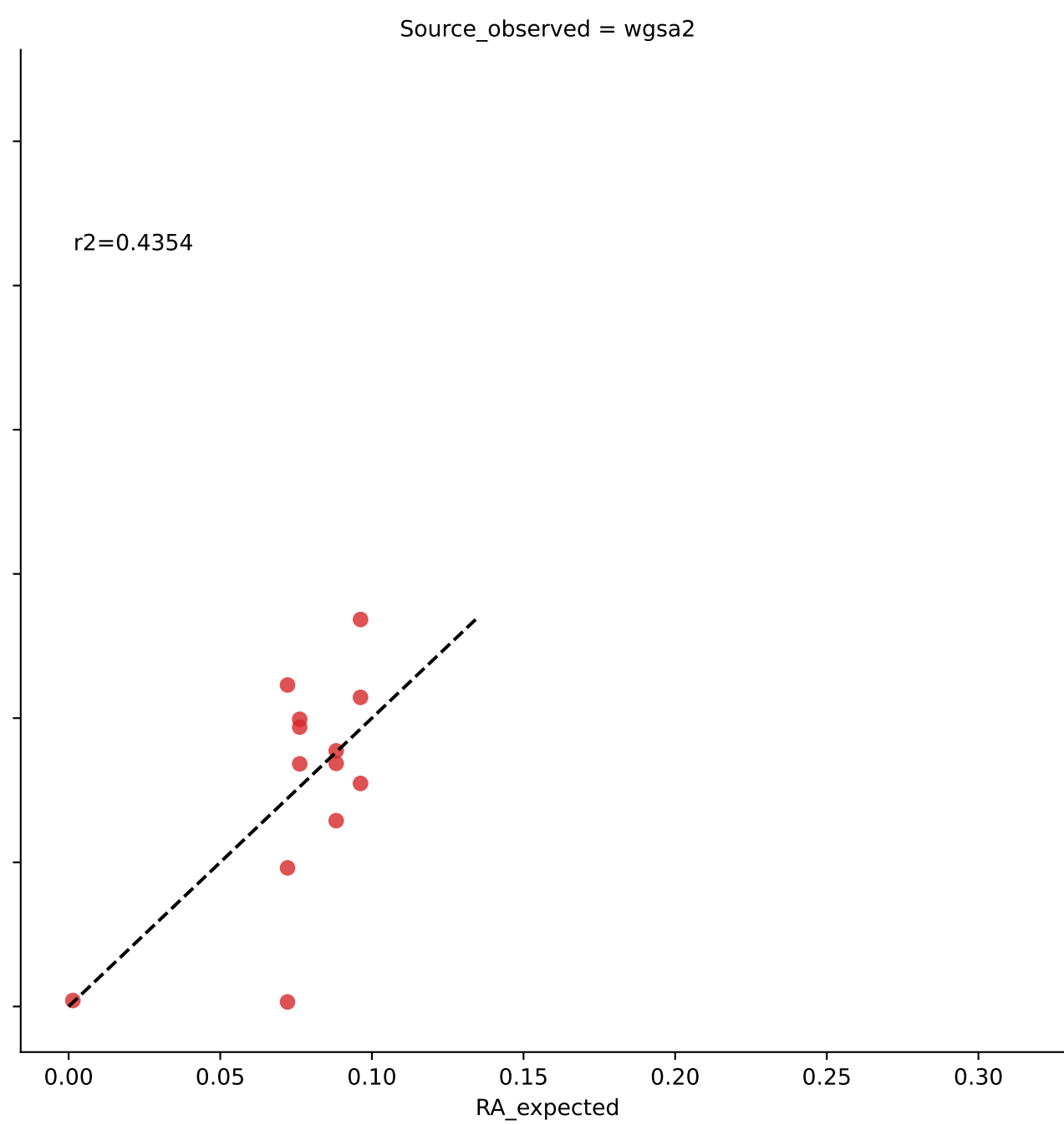
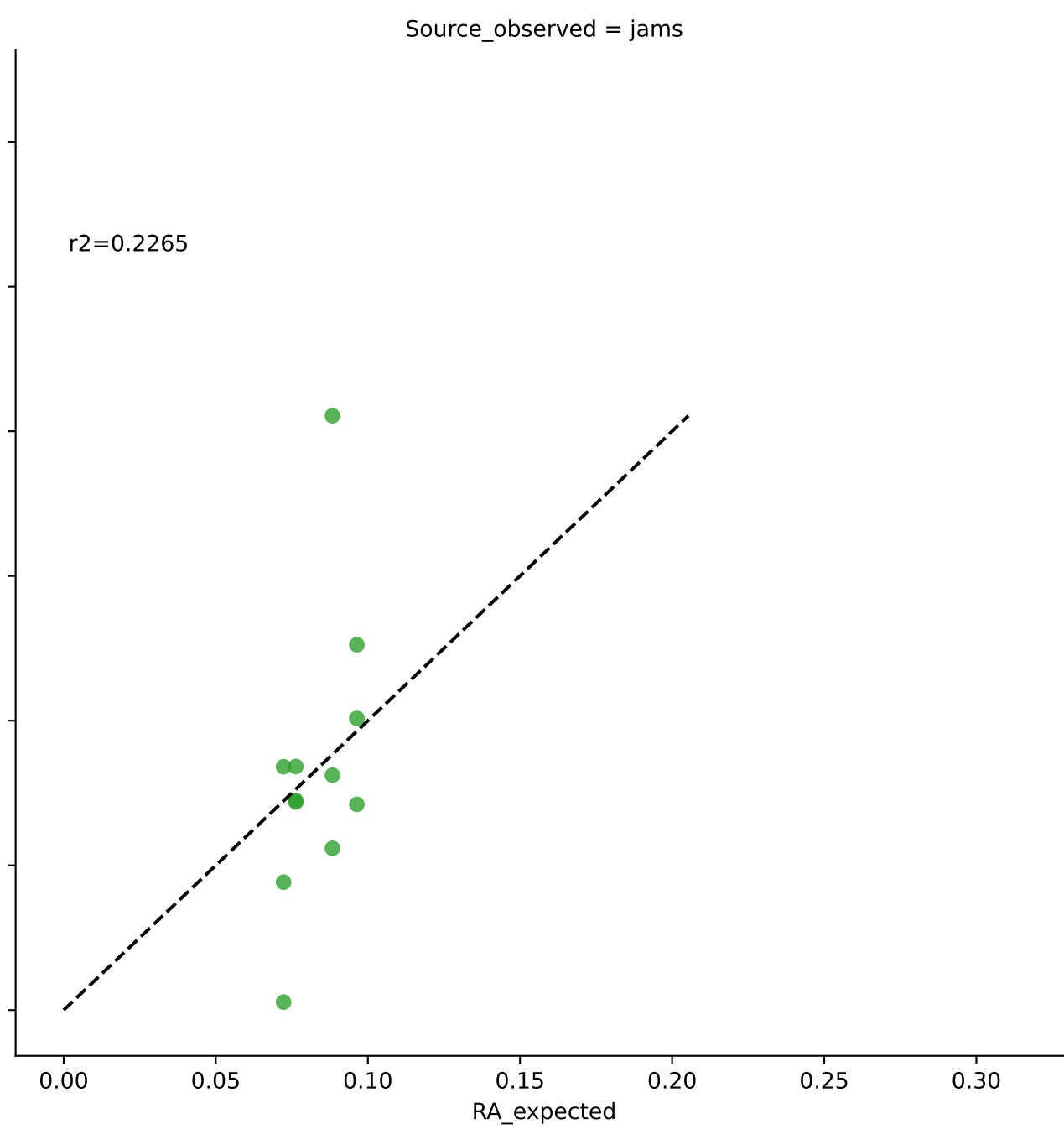
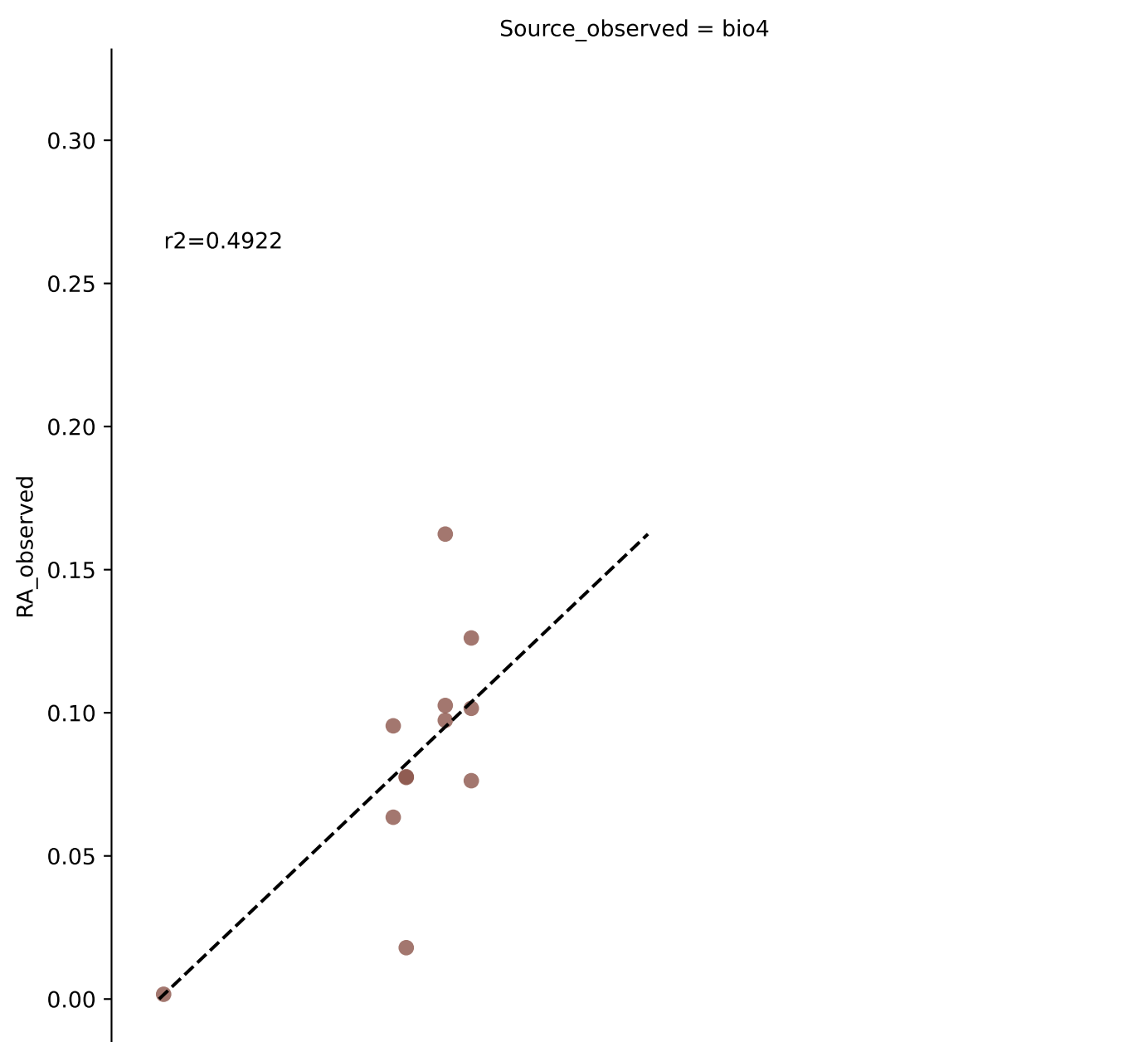


Bivariate Linear Regression for Sample S2 in Experiment camisimGI (Species at filter threshold 0.001)

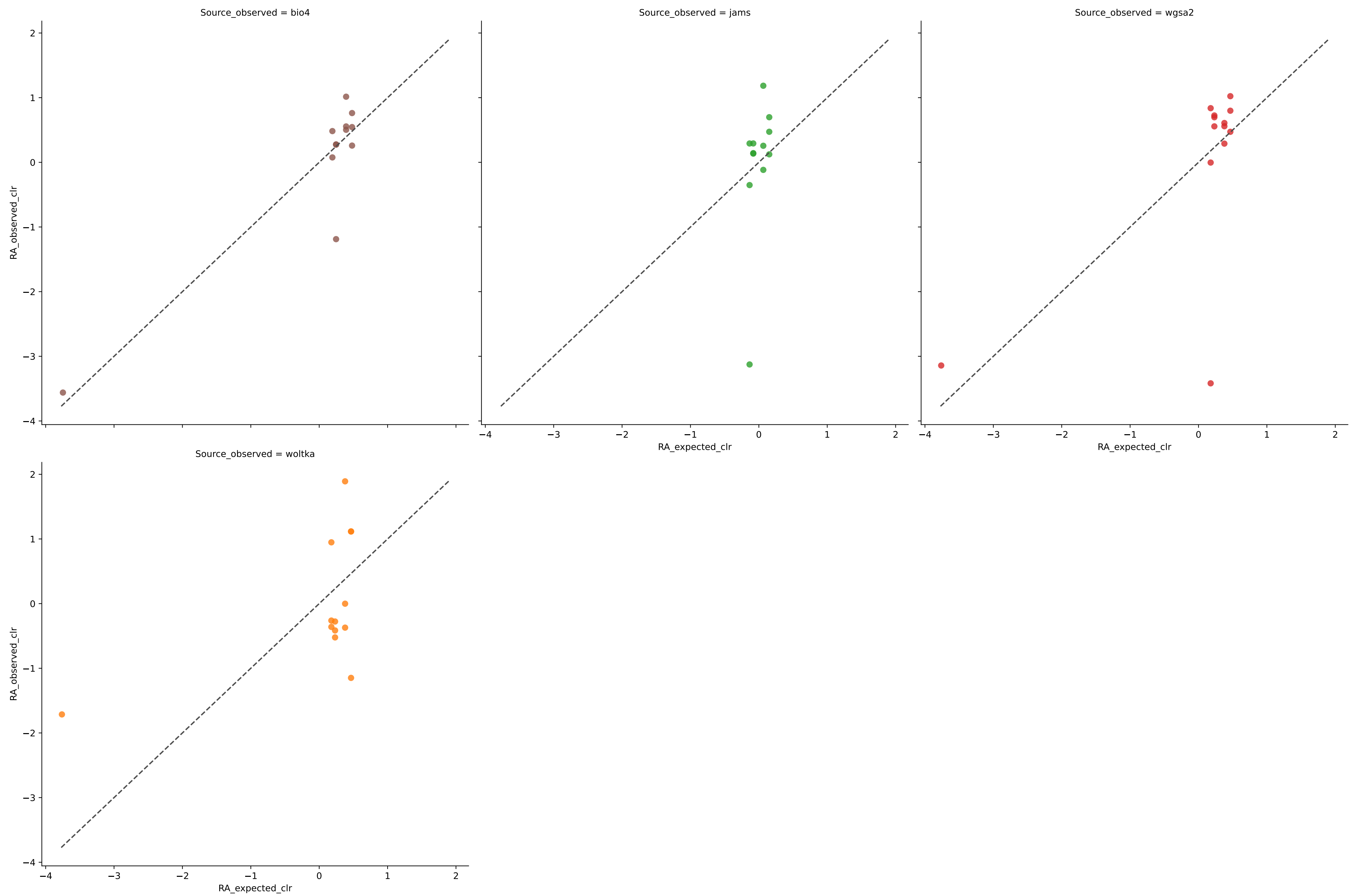


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	20	0.9598	0.0074	0.9077	0.9264	0.0195	100.0000	0.0000
jams	19	0.9386	0.0117	1.3696	0.8891	0.0213	100.0000	0.0000
wgsa	19	0.6551	0.0209	1.3118	0.8014	0.0504	100.0000	0.0000
woltka	16	0.9178	0.0192	1.7436	0.8467	0.0320	100.0000	0.0000

Bivariate Linear Regression for Sample EG in Experiment nist (Genus at filter threshold 0.001)

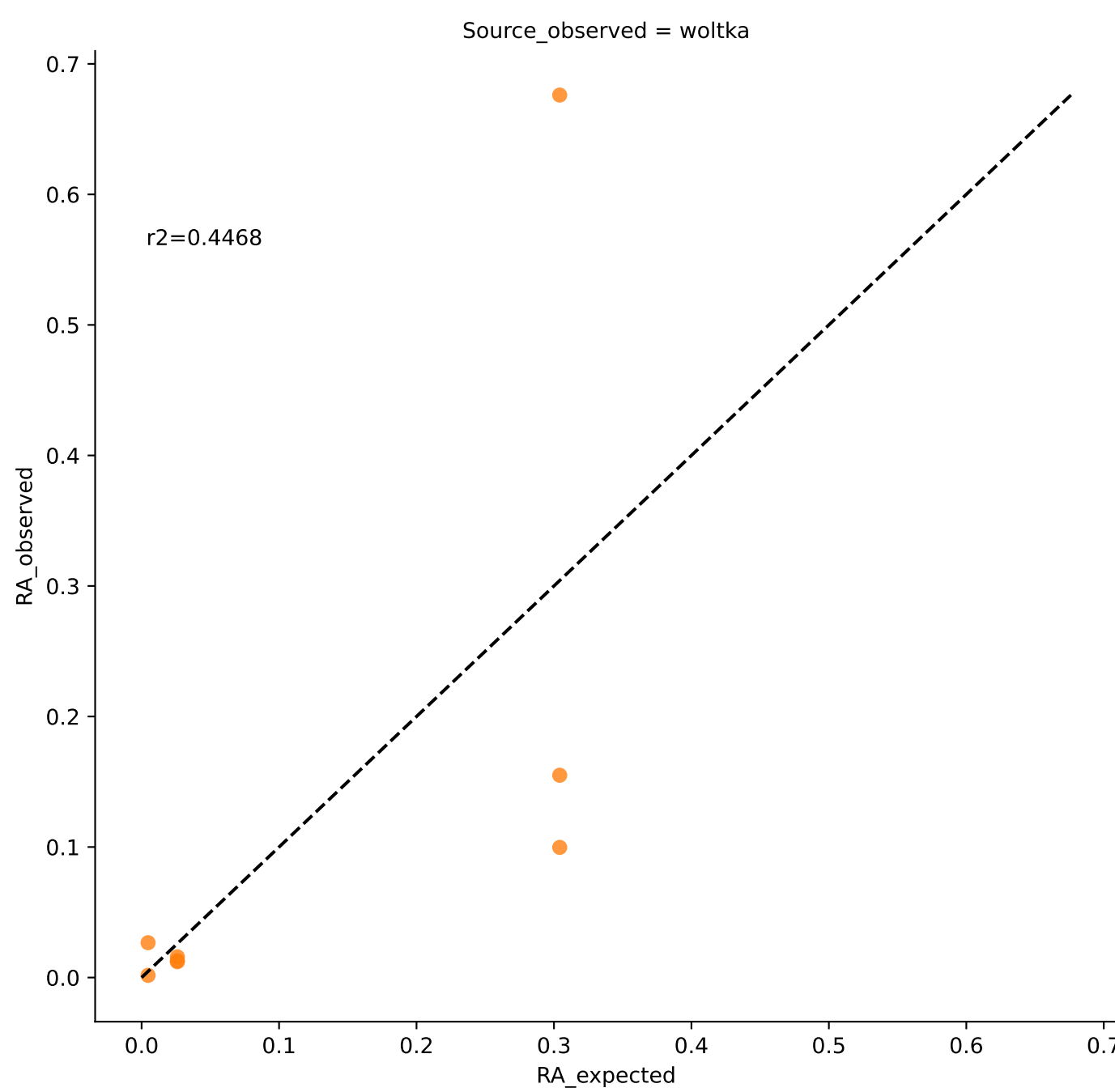
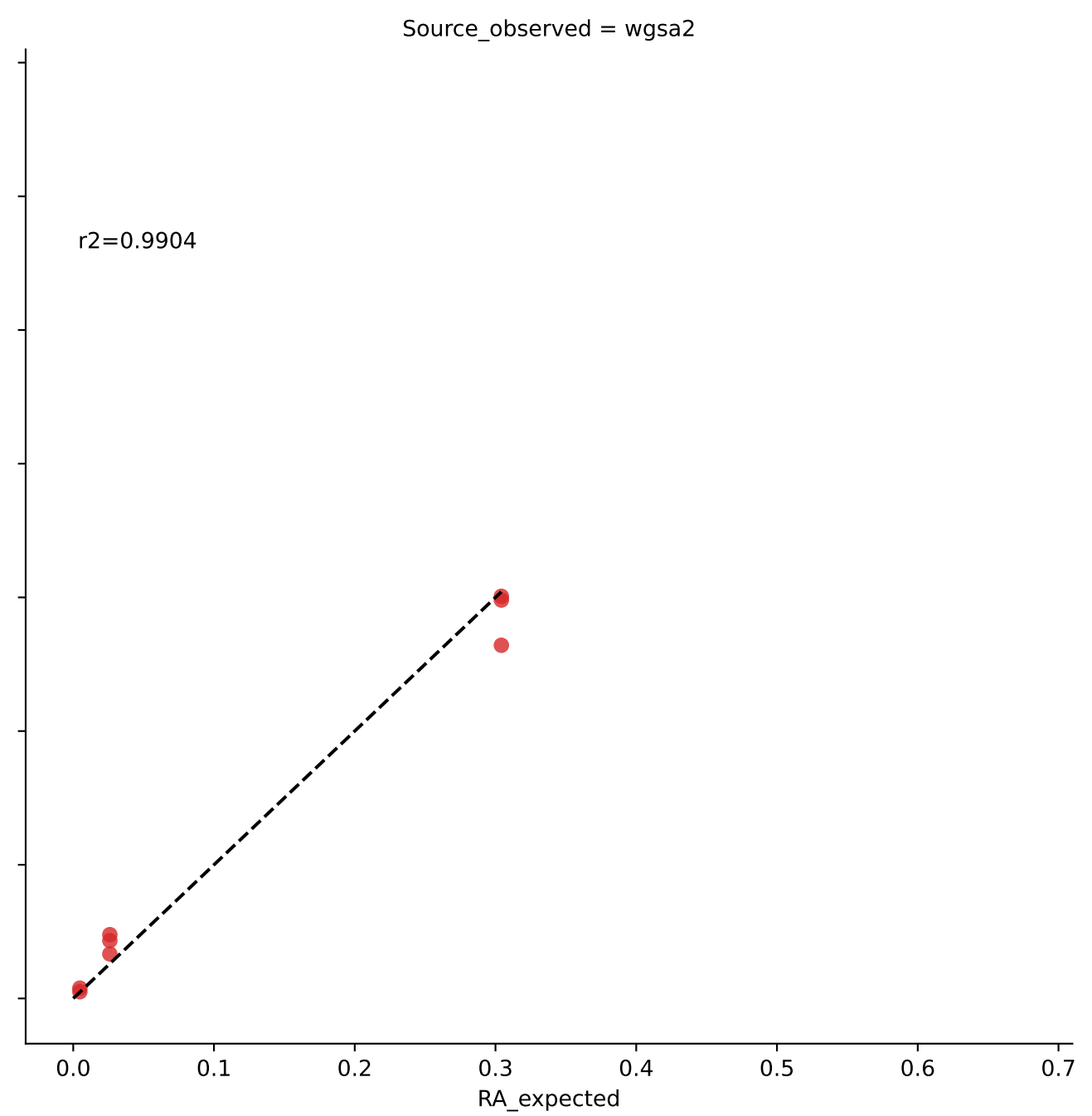
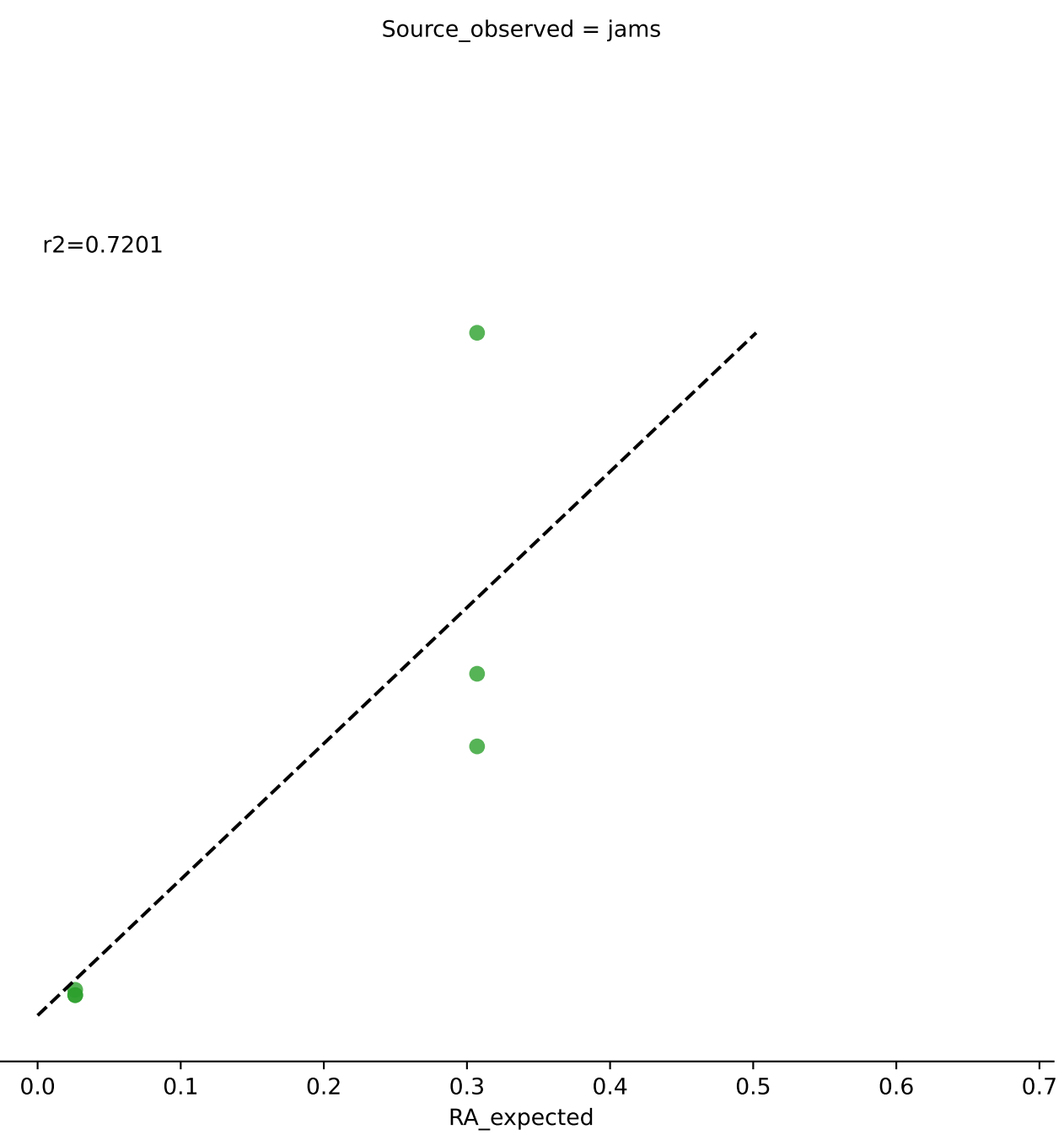
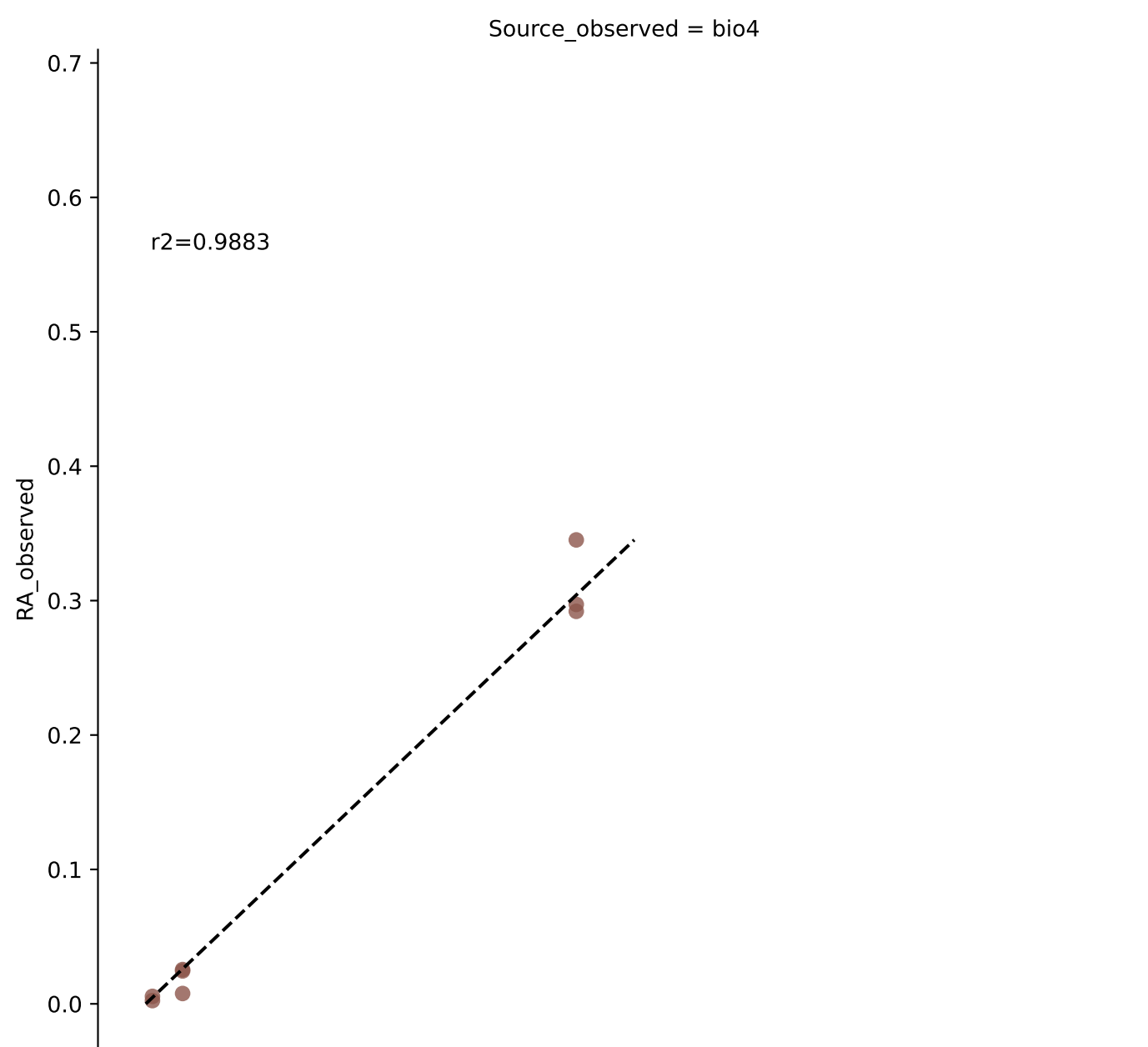


Bivariate Linear Regression for Sample EG in Experiment nist (Genus at filter threshold 0.001)

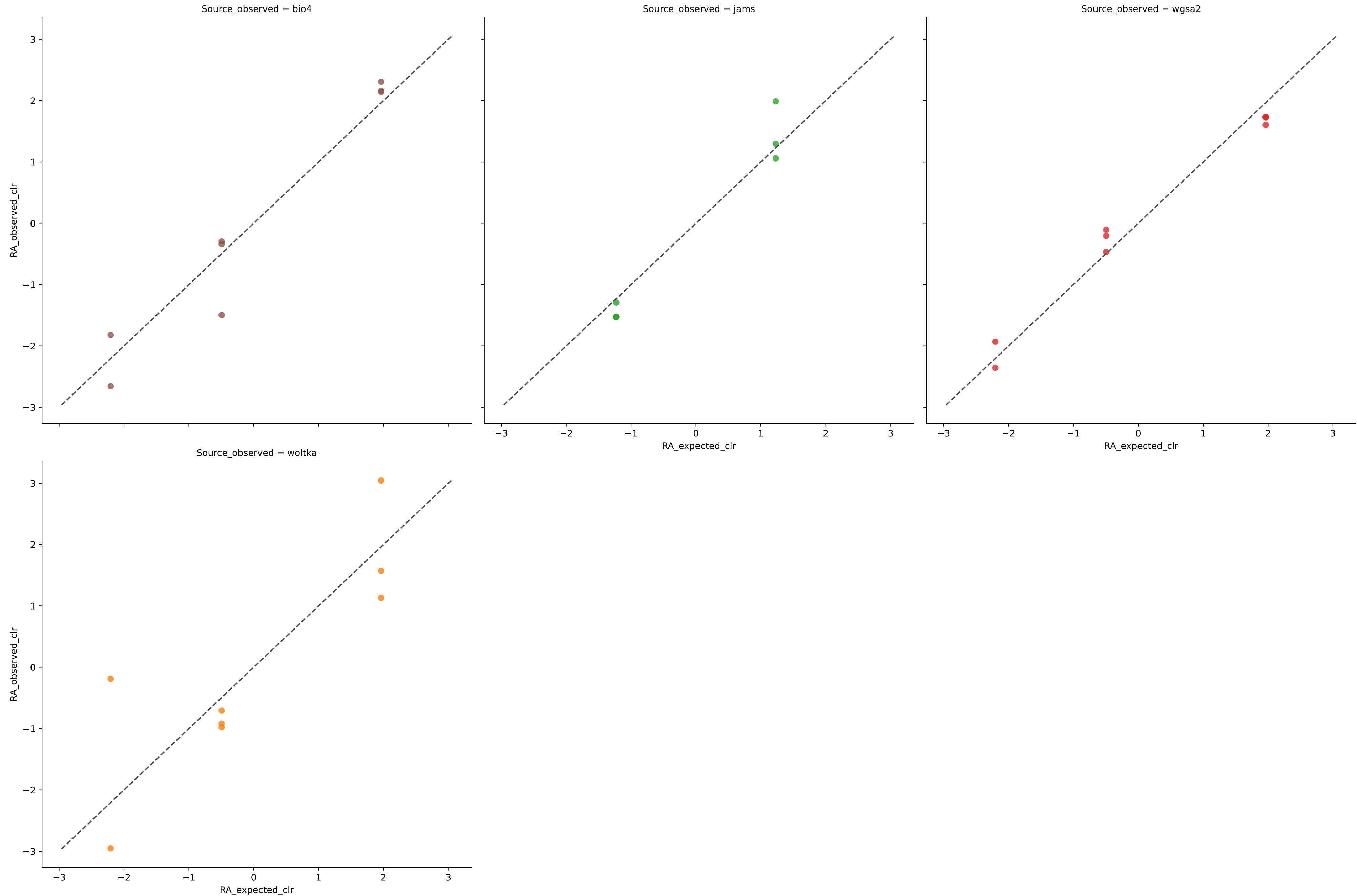


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	12	0.4922	0.0195	1.6584	0.8827	0.0296	100.0000	0.0000
jams	12	0.2265	0.0285	3.3366	0.8291	0.0430	100.0000	0.0000
wgsa2	13	0.4354	0.0217	3.8549	0.8587	0.0286	100.0000	0.0000
woltka	13	0.1179	0.0591	3.6010	0.6157	0.0781	100.0000	0.0000

Bivariate Linear Regression for Sample MIX-A in Experiment nist (Genus at filter threshold 0.001)

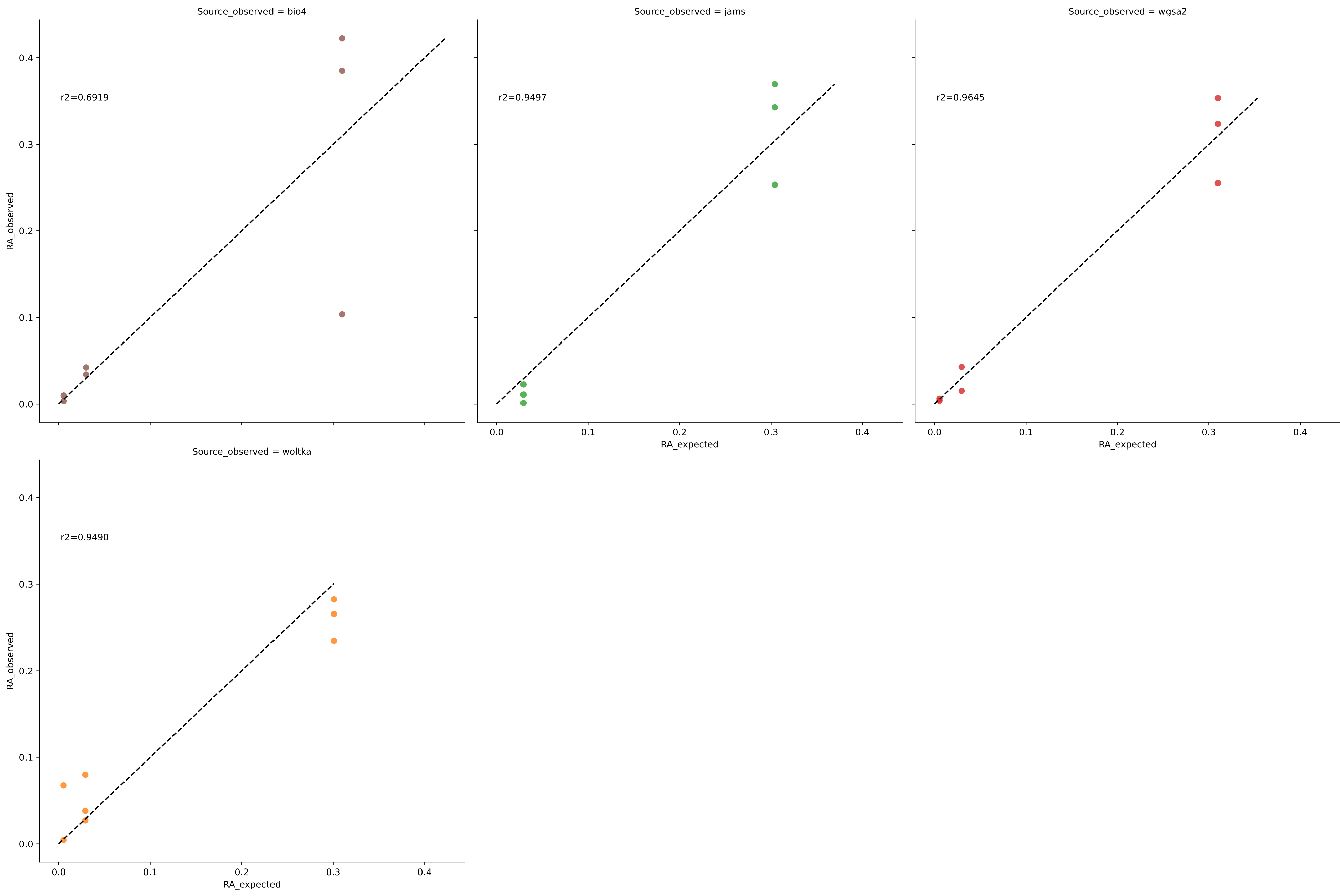


Bivariate Linear Regression for Sample MIX-A in Experiment nist (Genus at filter threshold 0.001)

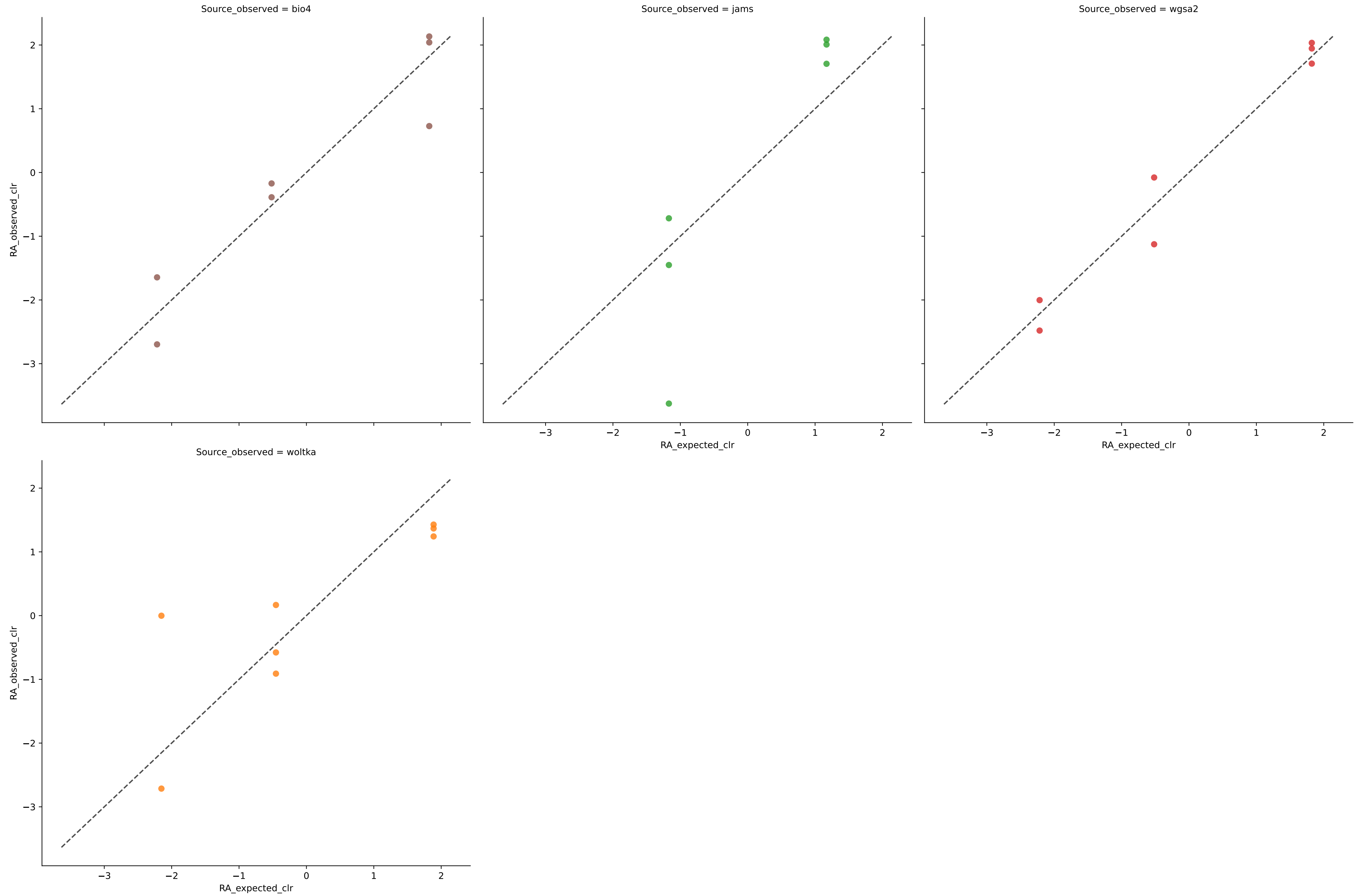


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	8	0.9883	0.0105	1.2667	0.9582	0.0167	100.0000	0.0000
jams	6	0.7201	0.0650	0.8890	0.8050	0.0943	100.0000	0.0000
wgsa2	8	0.9904	0.0124	0.7571	0.9505	0.0176	100.0000	0.0000
woltka	8	0.4468	0.0985	2.6646	0.6061	0.1594	100.0000	0.0000

Bivariate Linear Regression for Sample MIX-B in Experiment nist (Genus at filter threshold 0.001)

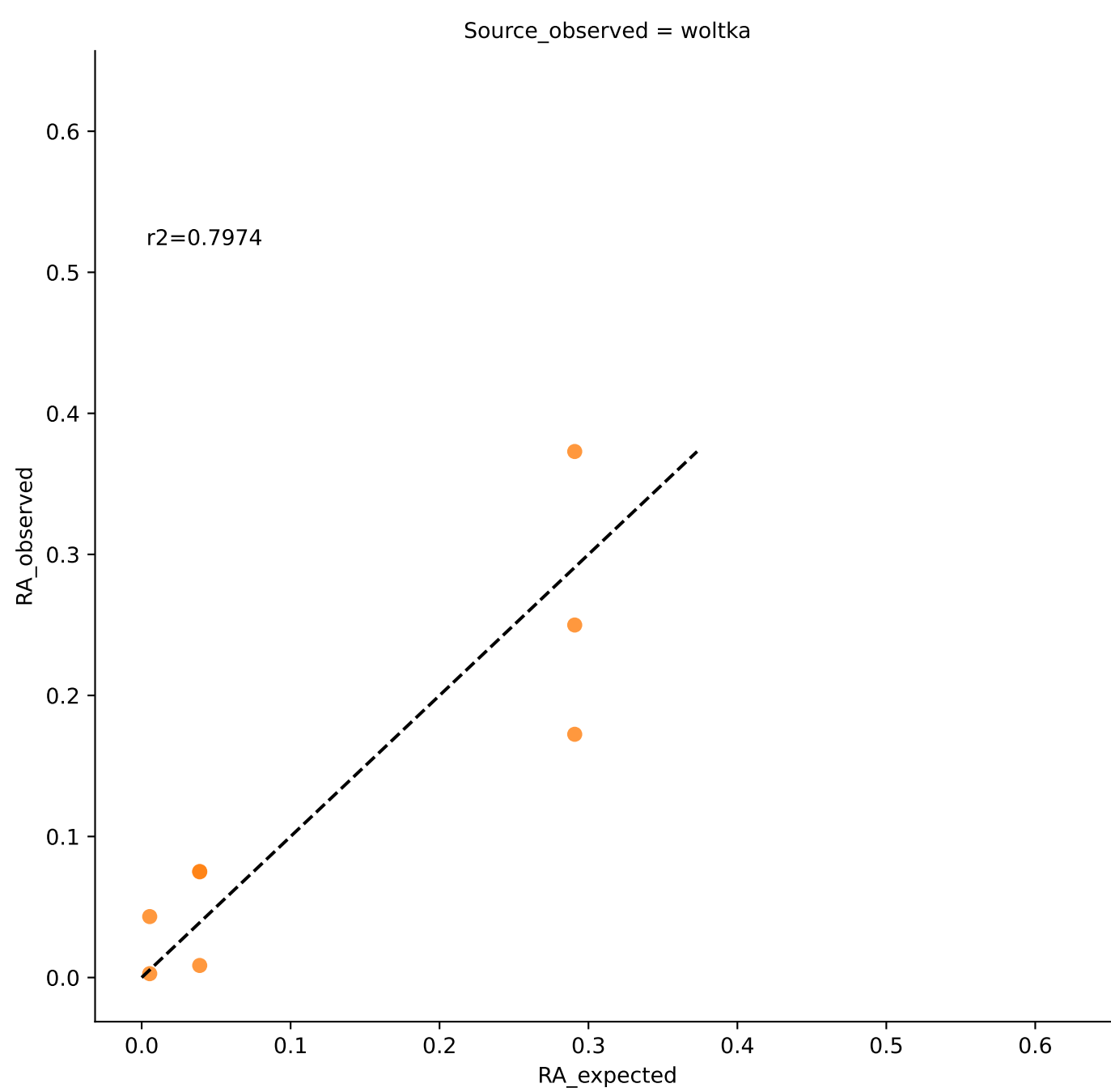
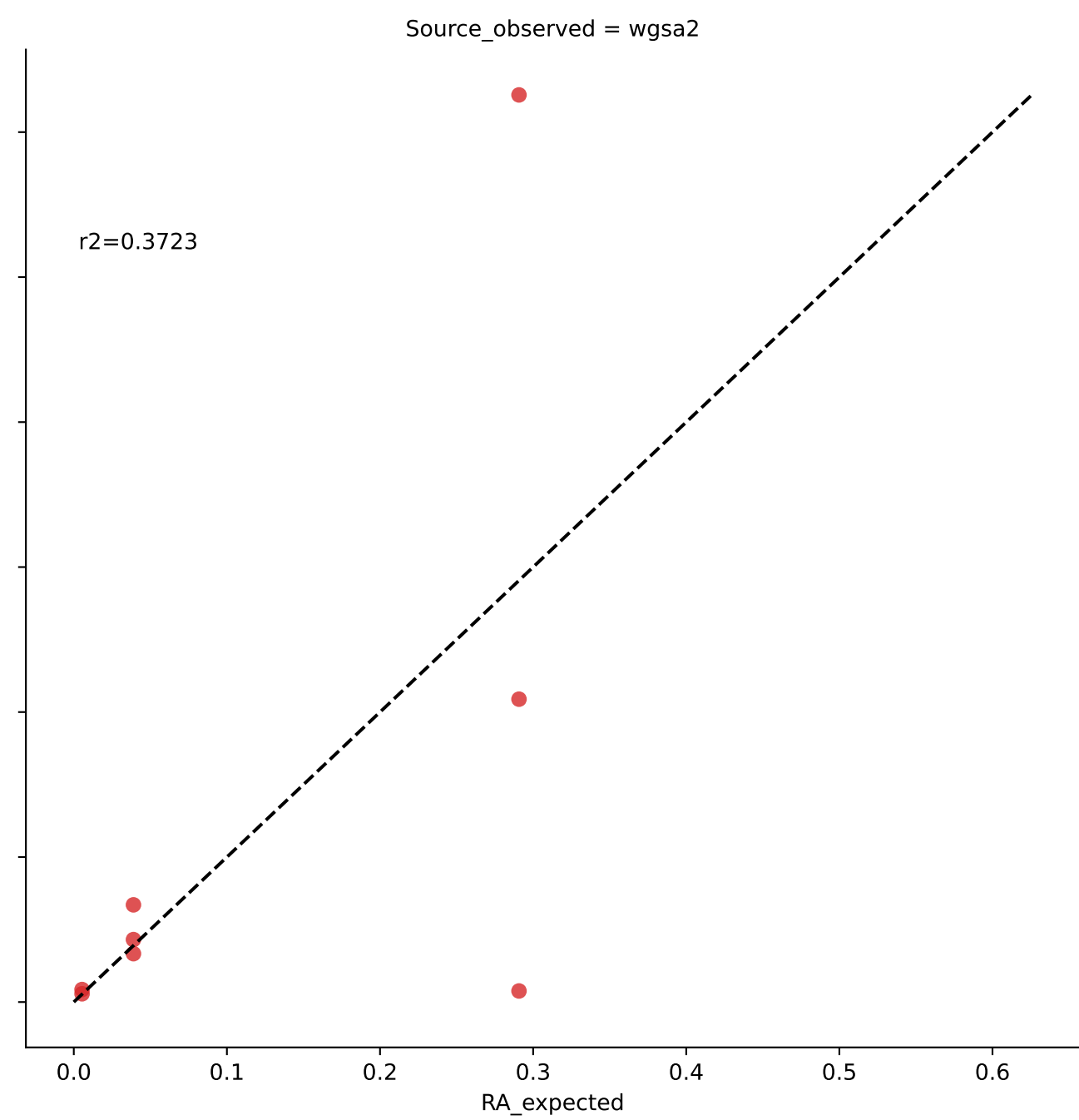
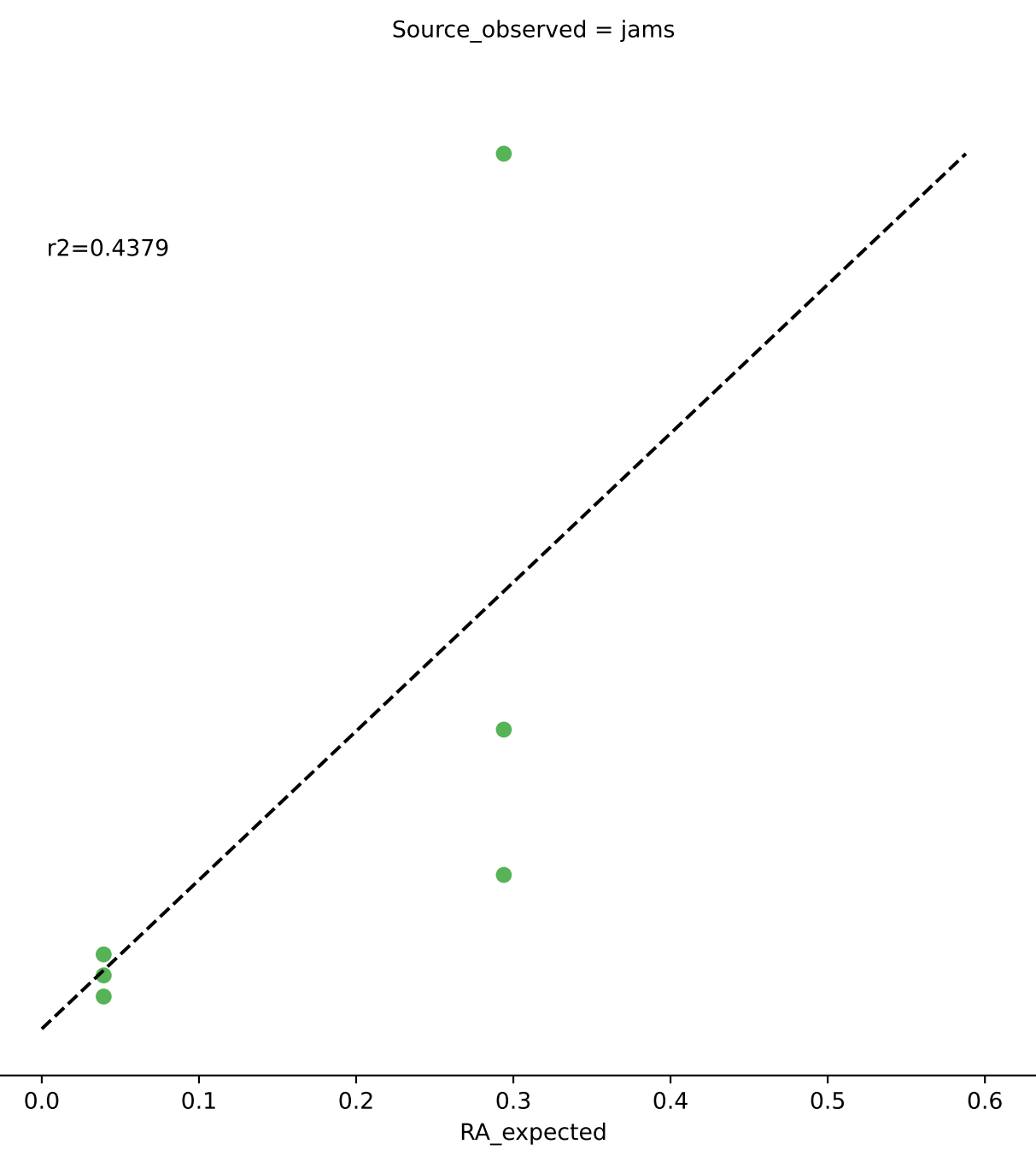
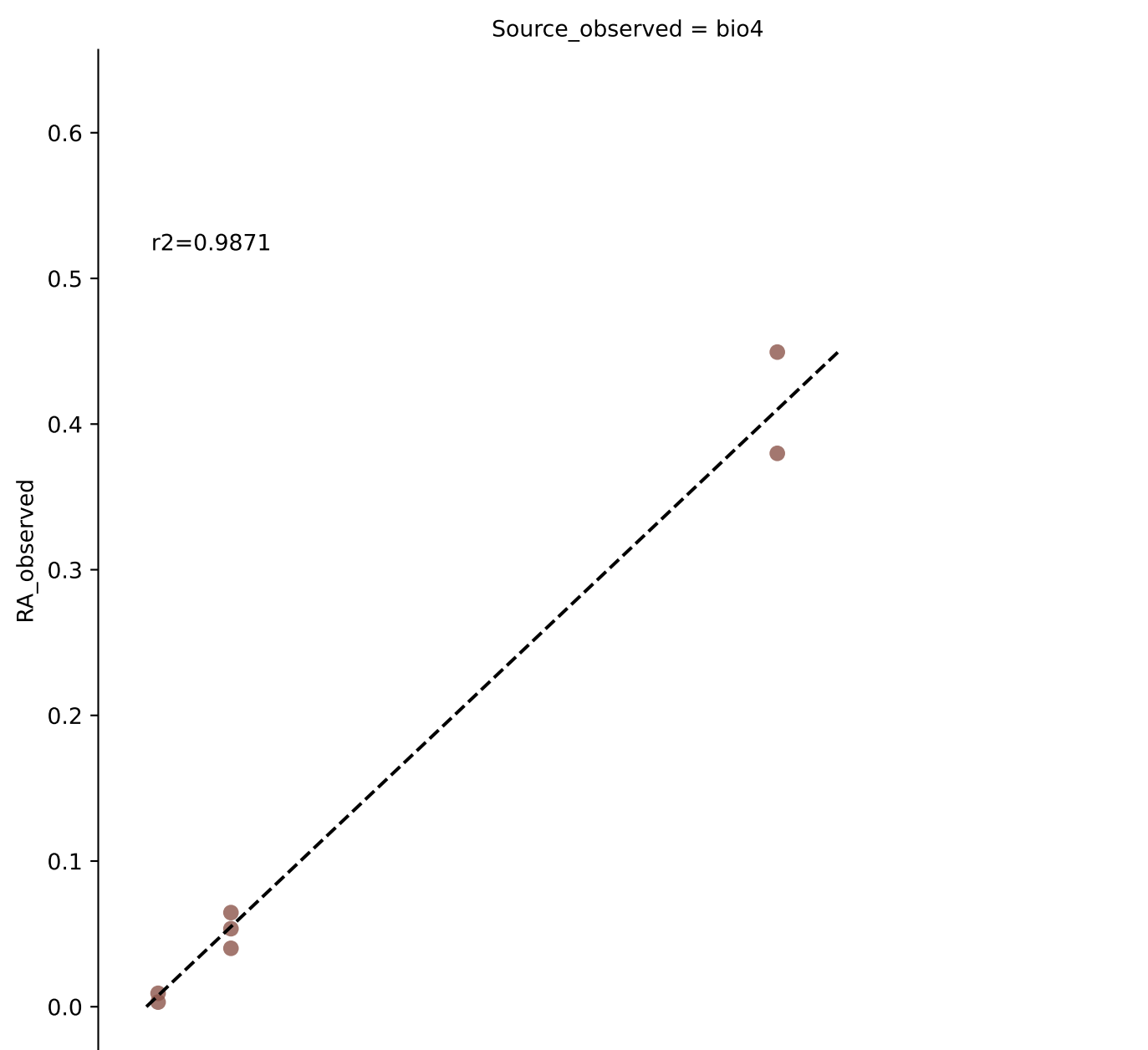


Bivariate Linear Regression for Sample MIX-B in Experiment nist (Genus at filter threshold 0.001)

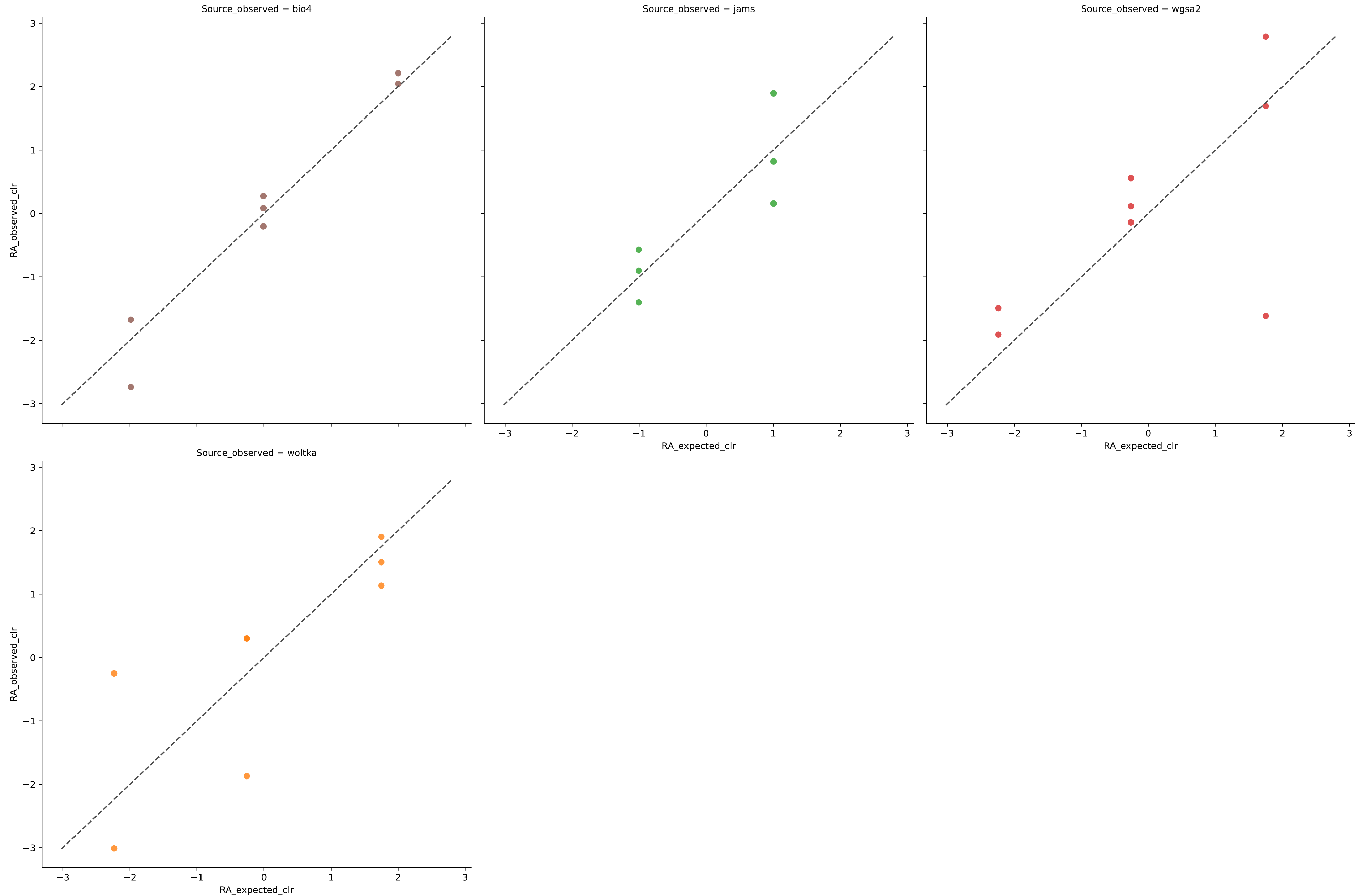


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	7	0.6919	0.0595	1.4266	0.7917	0.0934	100.0000	0.0000
jams	6	0.9497	0.0347	2.8524	0.8958	0.0399	100.0000	0.0000
wgsa2	7	0.9645	0.0203	0.8667	0.9290	0.0279	100.0000	0.0000
woltka	8	0.9490	0.0306	2.5389	0.8777	0.0396	100.0000	0.0000

Bivariate Linear Regression for Sample MIX-C in Experiment nist (Genus at filter threshold 0.001)

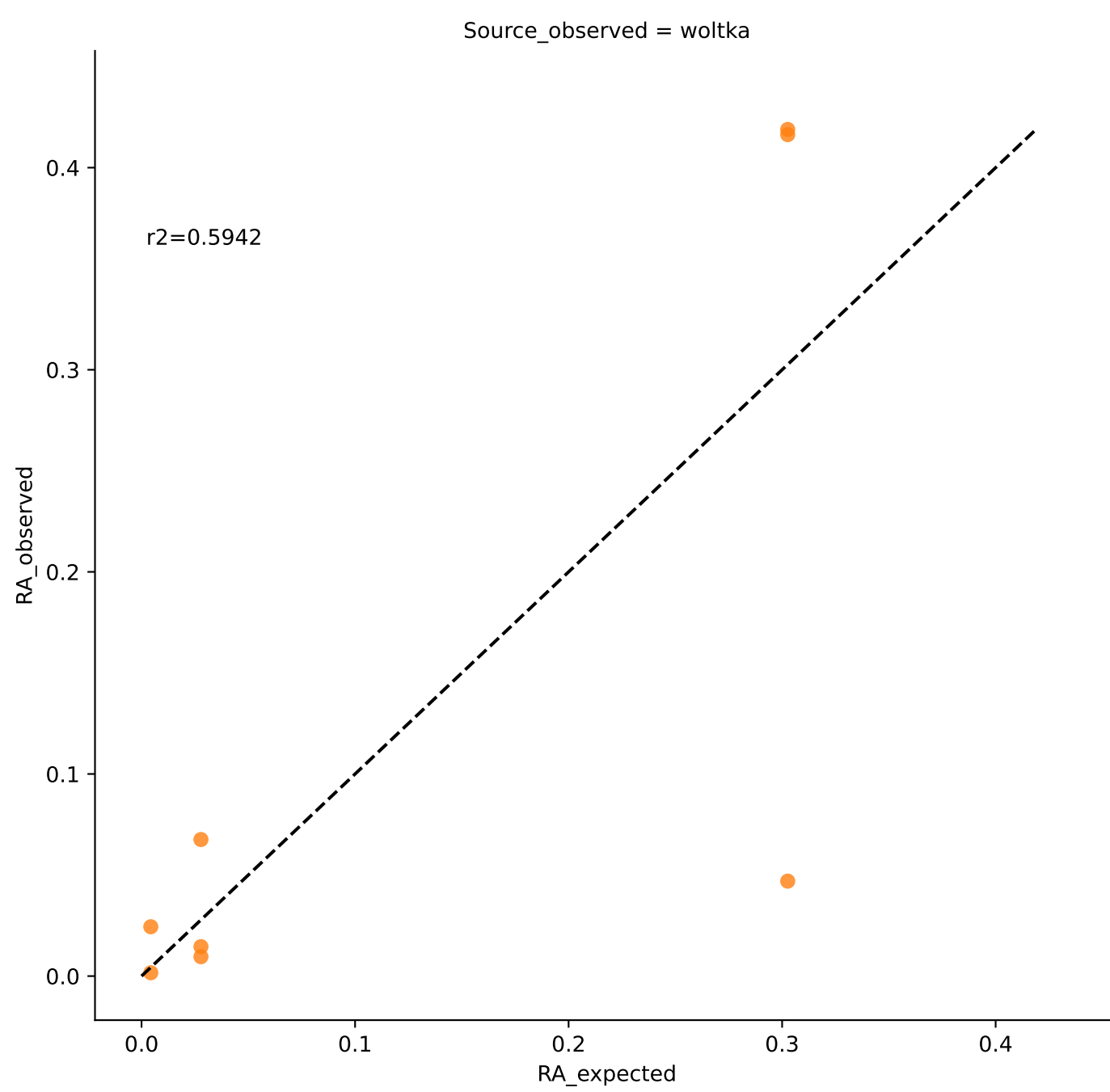
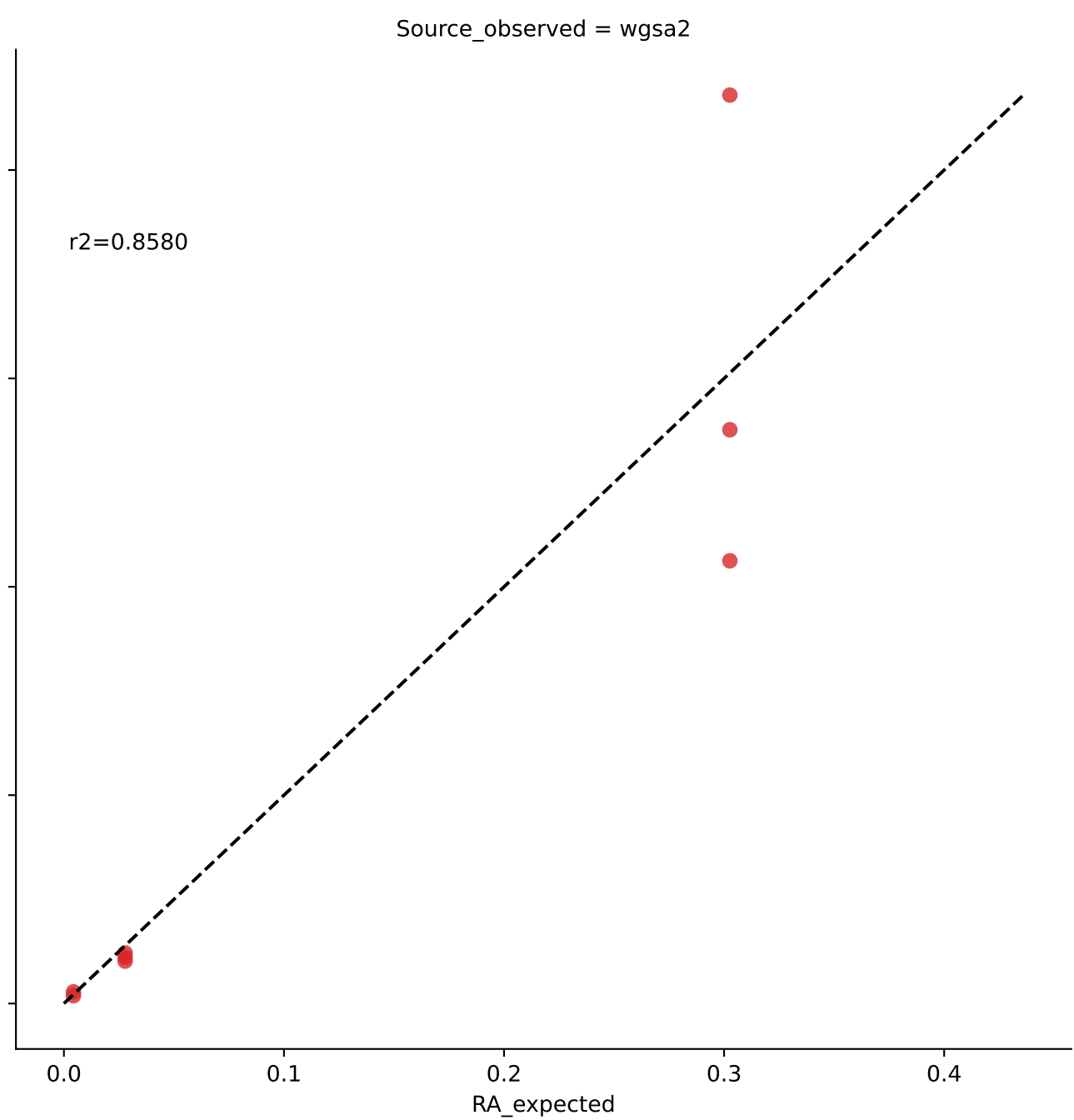
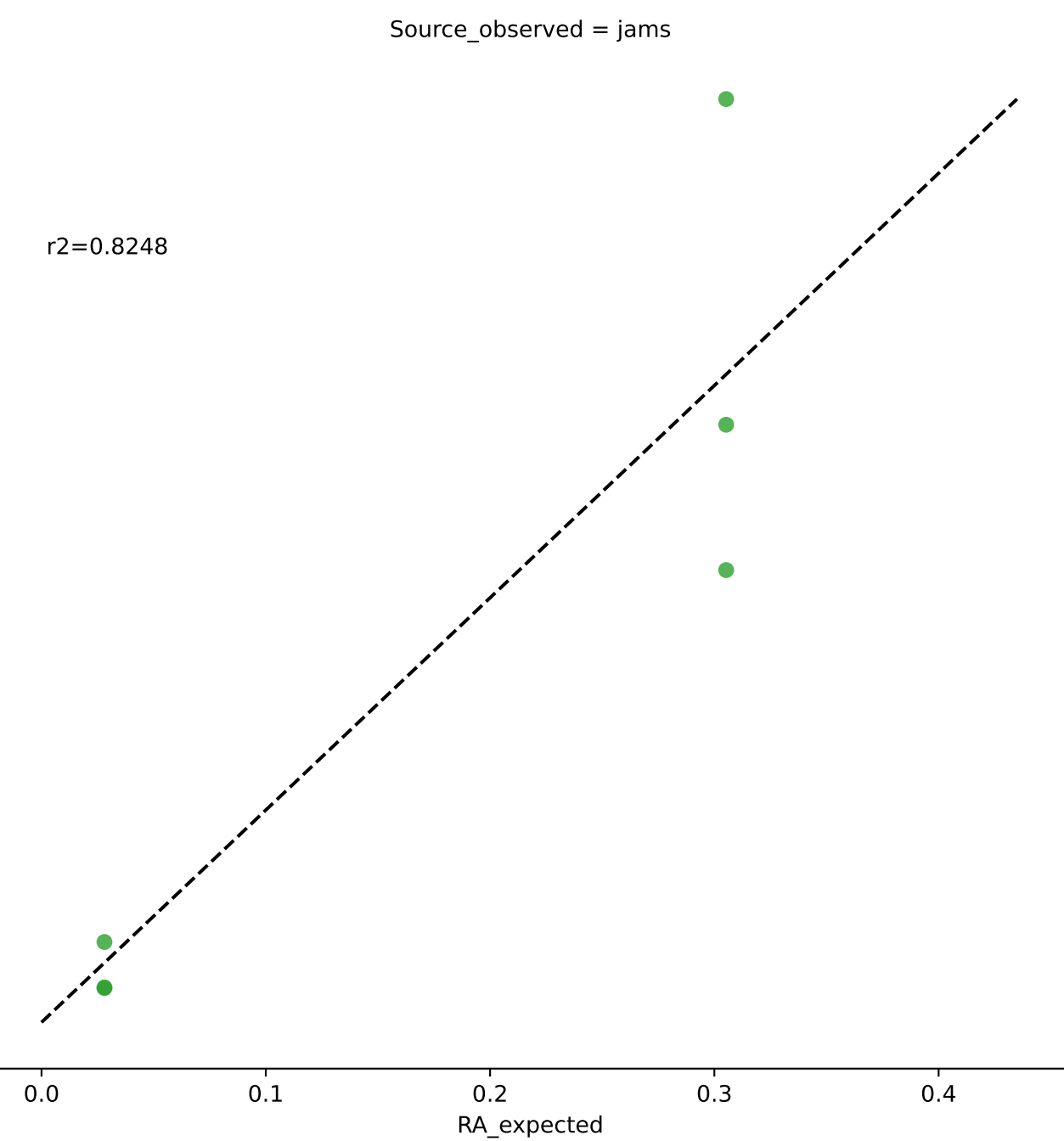
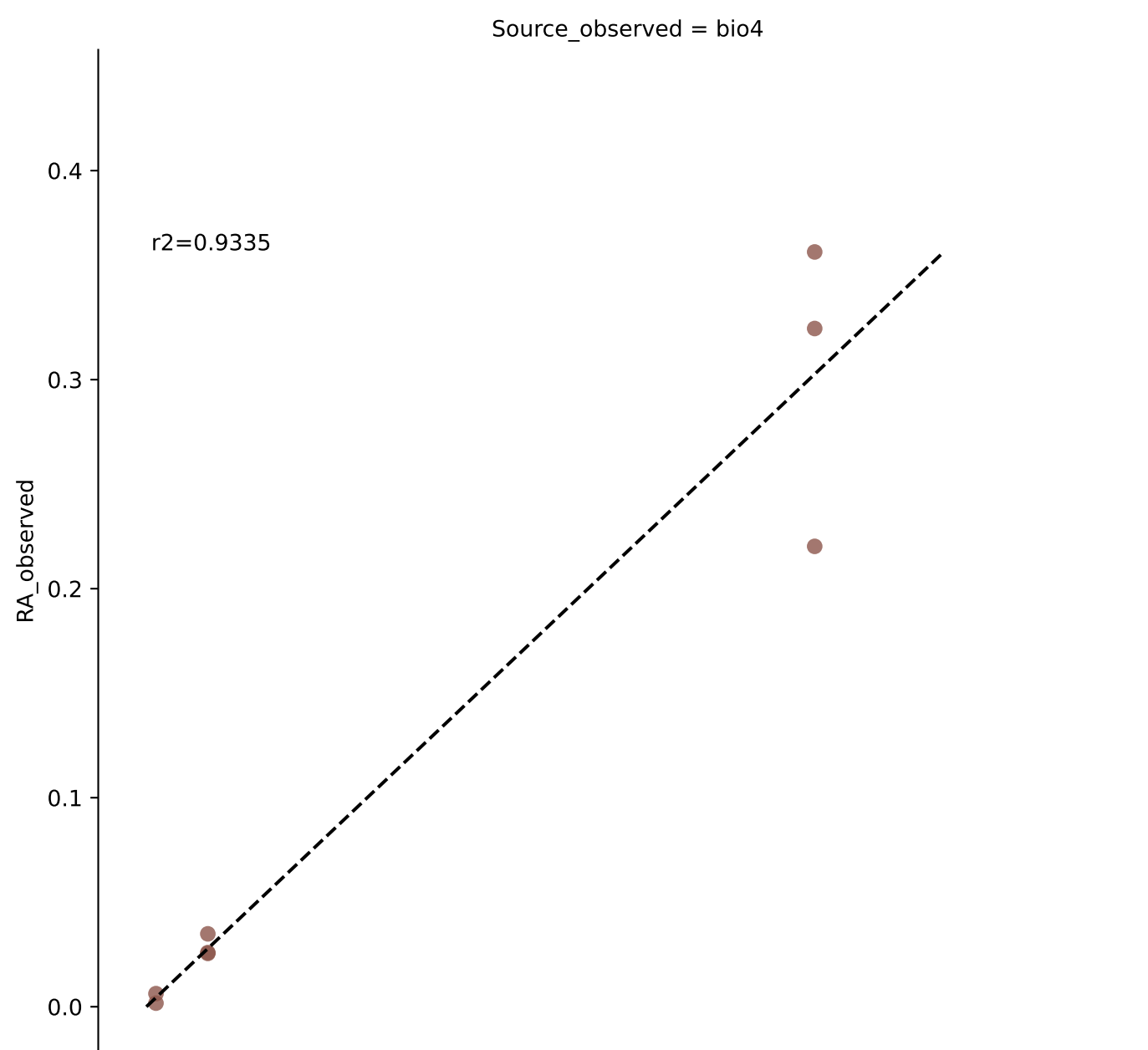


Bivariate Linear Regression for Sample MIX-C in Experiment nist (Genus at filter threshold 0.001)

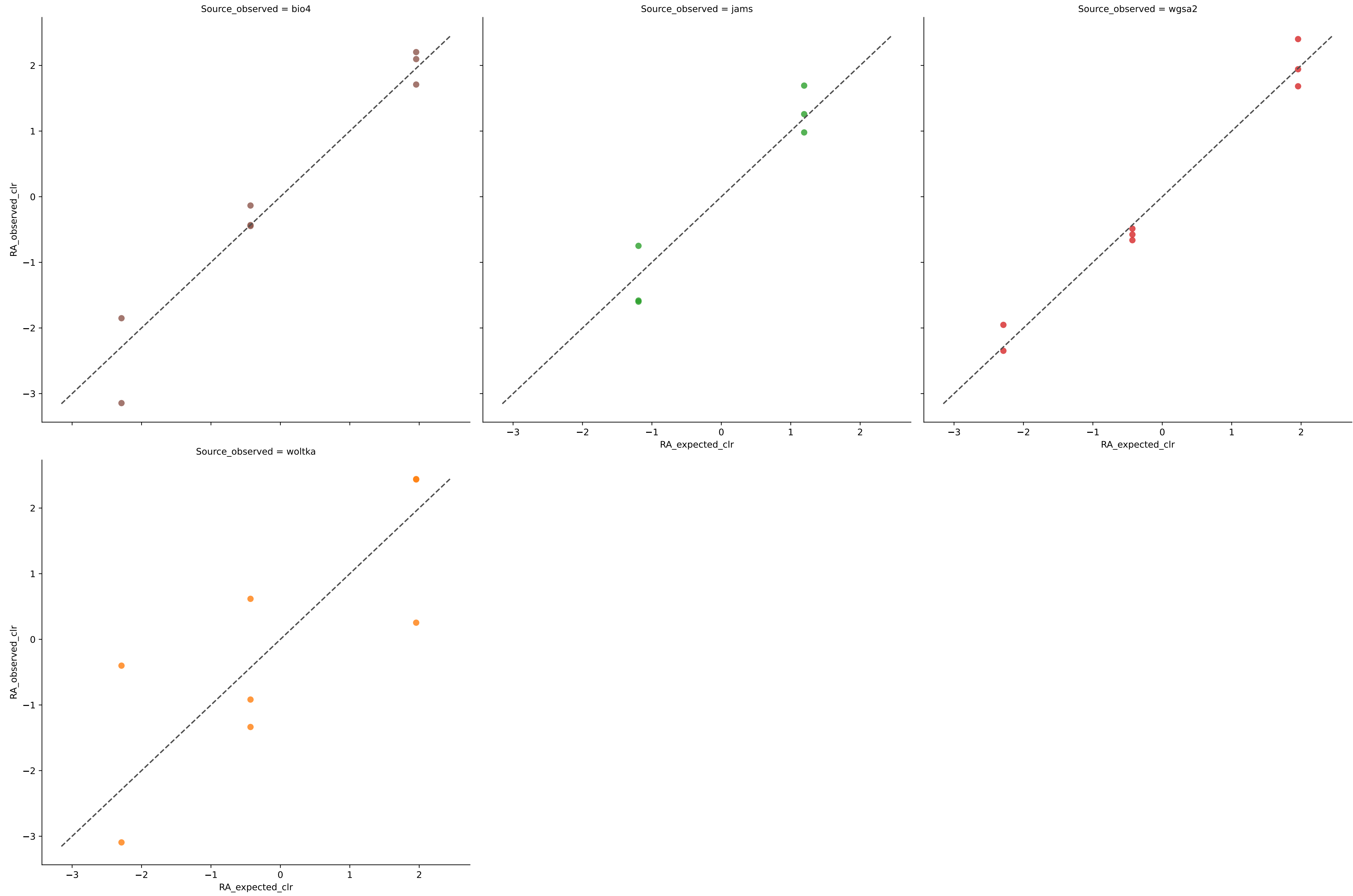


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	7	0.9871	0.0145	0.9149	0.9493	0.0200	100.0000	0.0000
jams	6	0.4379	0.1015	1.3797	0.6954	0.1482	100.0000	0.0000
wgsa2	8	0.3723	0.0927	3.7273	0.6294	0.1581	100.0000	0.0000
woltka	8	0.7974	0.0481	2.8685	0.8078	0.0585	100.0000	0.0000

Bivariate Linear Regression for Sample MIX-D in Experiment nist (Genus at filter threshold 0.001)

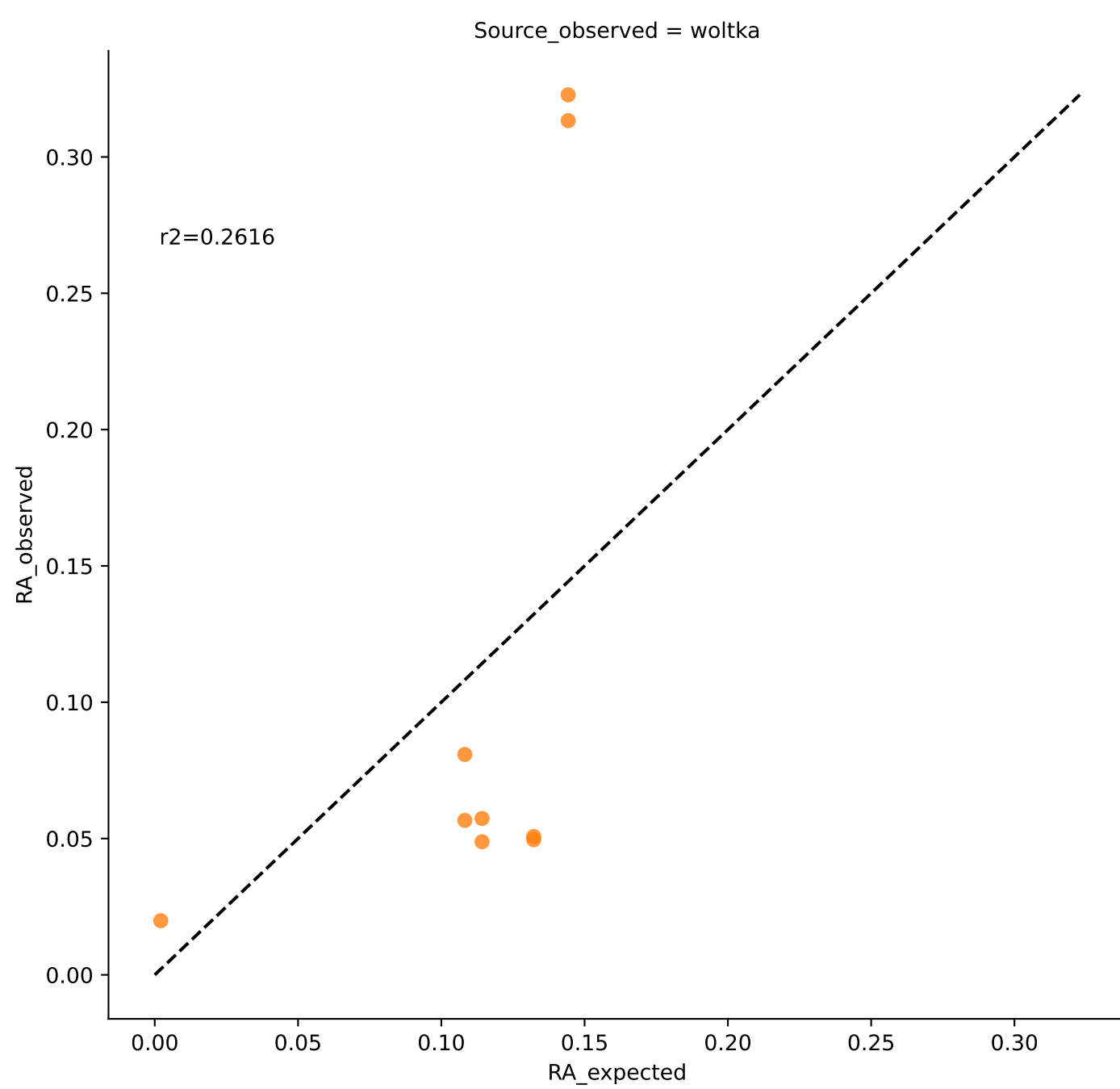
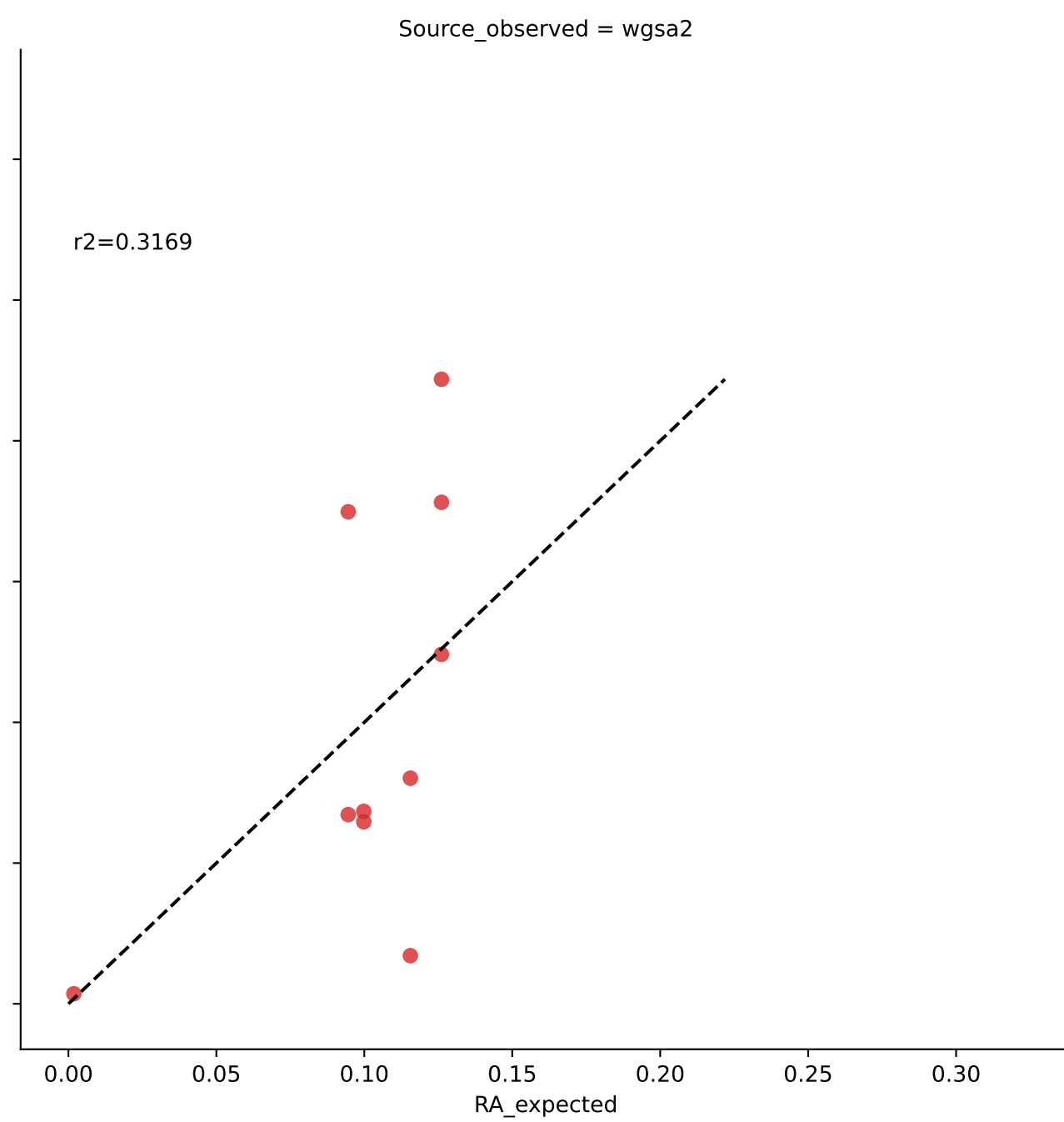
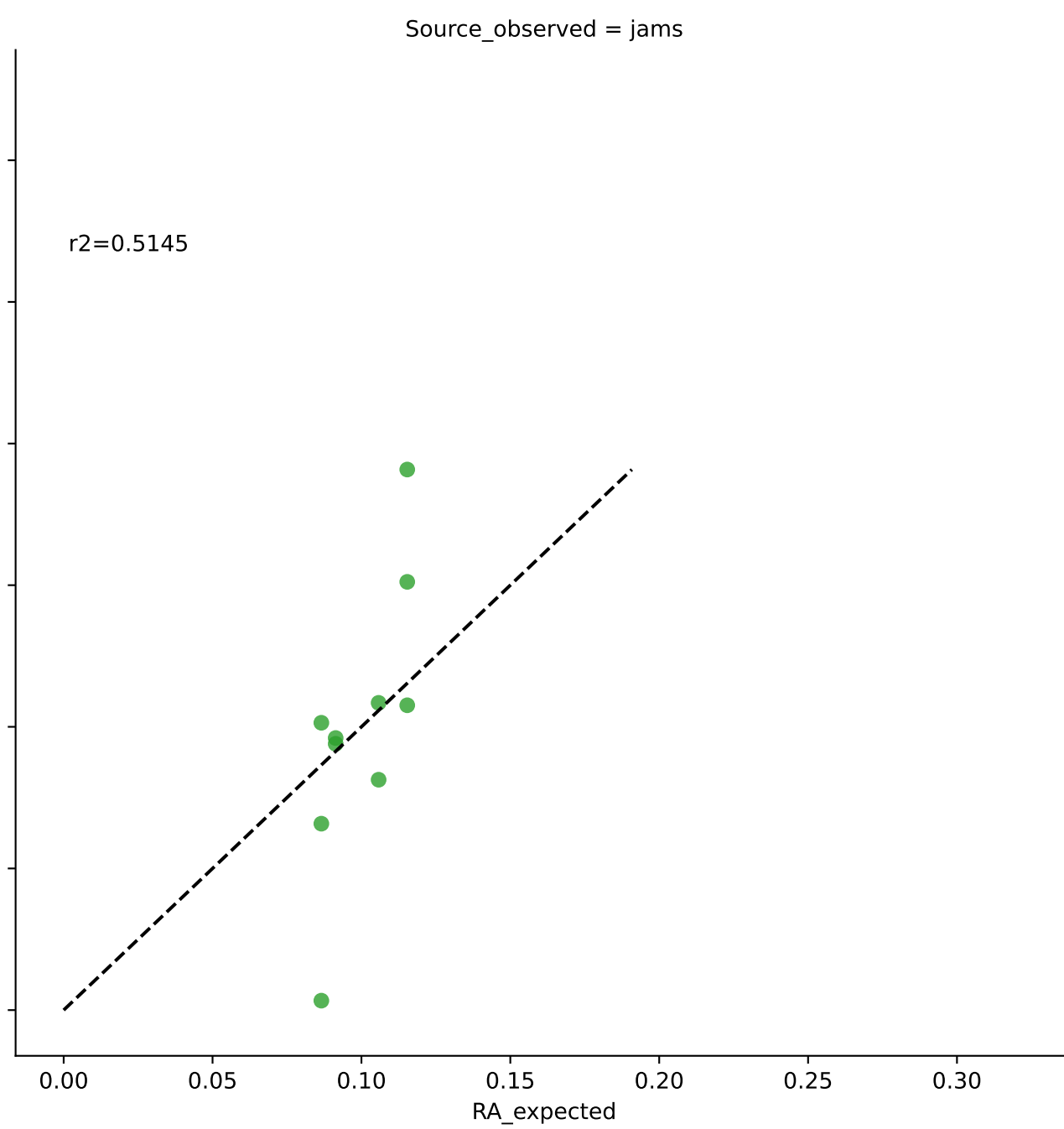
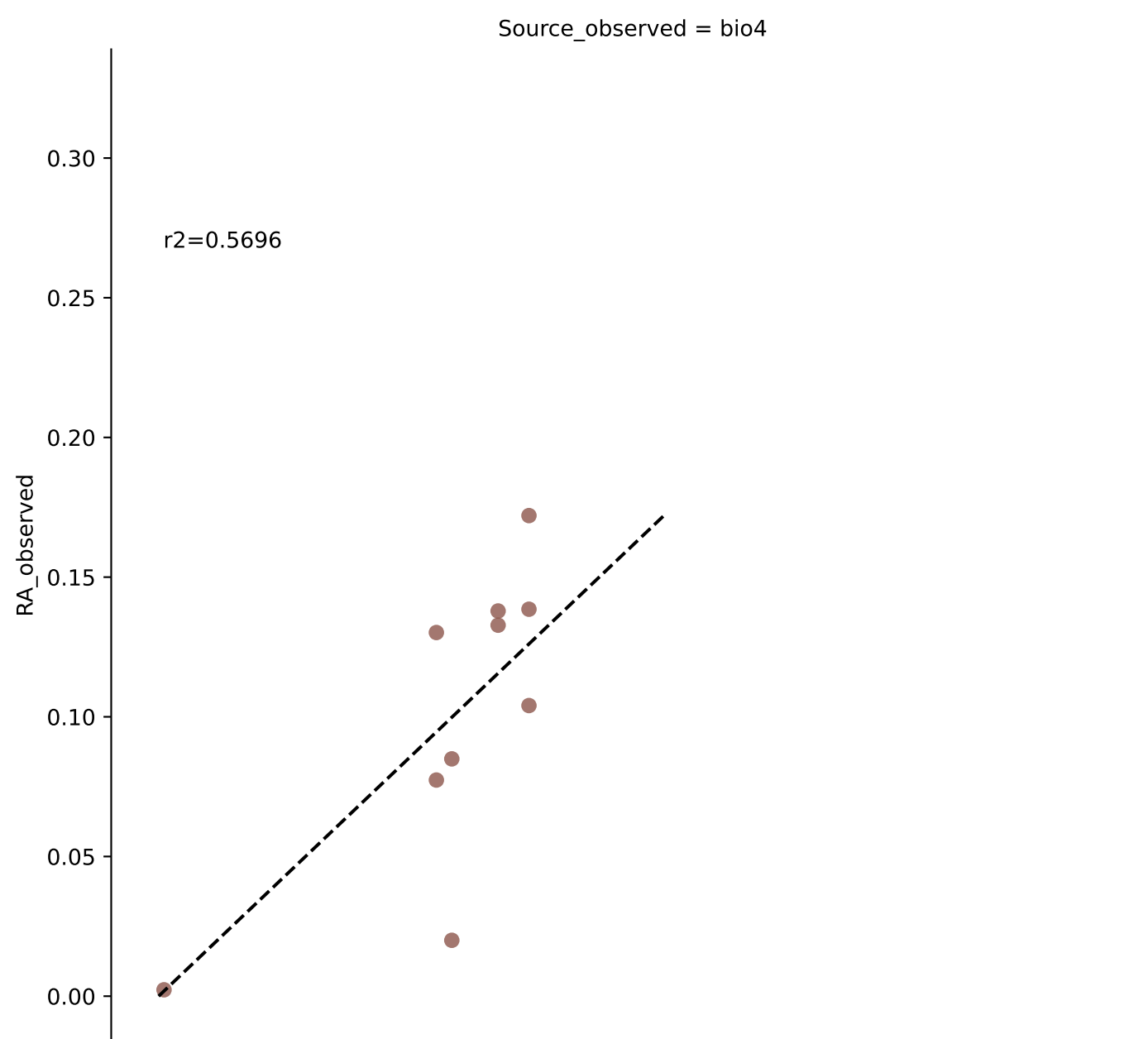


Bivariate Linear Regression for Sample MIX-D in Experiment nist (Genus at filter threshold 0.001)

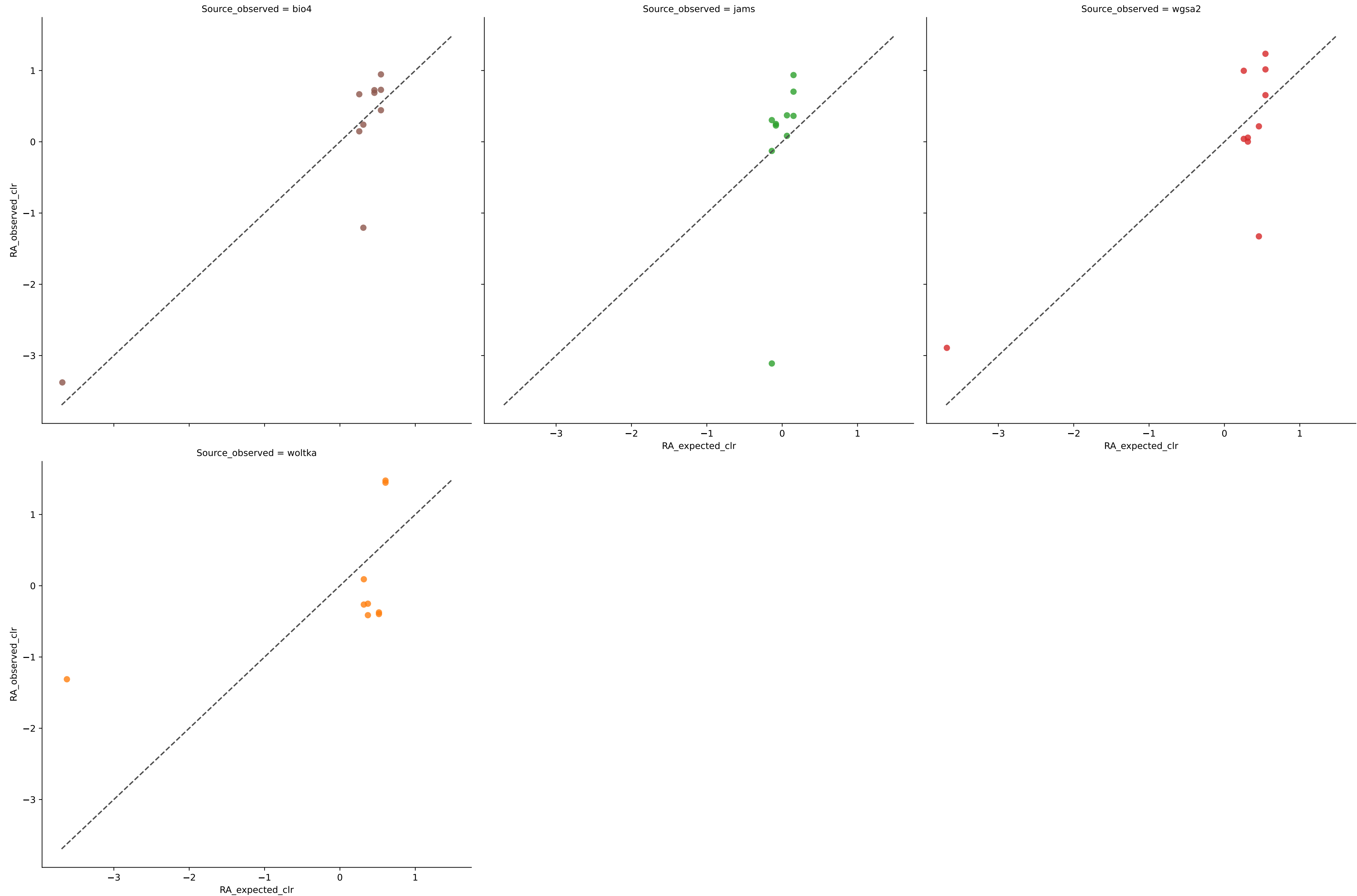


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	8	0.9335	0.0223	1.0739	0.9107	0.0367	100.0000	0.0000
jams	6	0.8248	0.0465	0.9017	0.8605	0.0661	100.0000	0.0000
wgsa2	8	0.8580	0.0337	0.6842	0.8654	0.0578	100.0000	0.0000
woltka	8	0.5942	0.0725	3.1193	0.7102	0.1086	100.0000	0.0000

Bivariate Linear Regression for Sample EG in Experiment nist (Species at filter threshold 0.001)

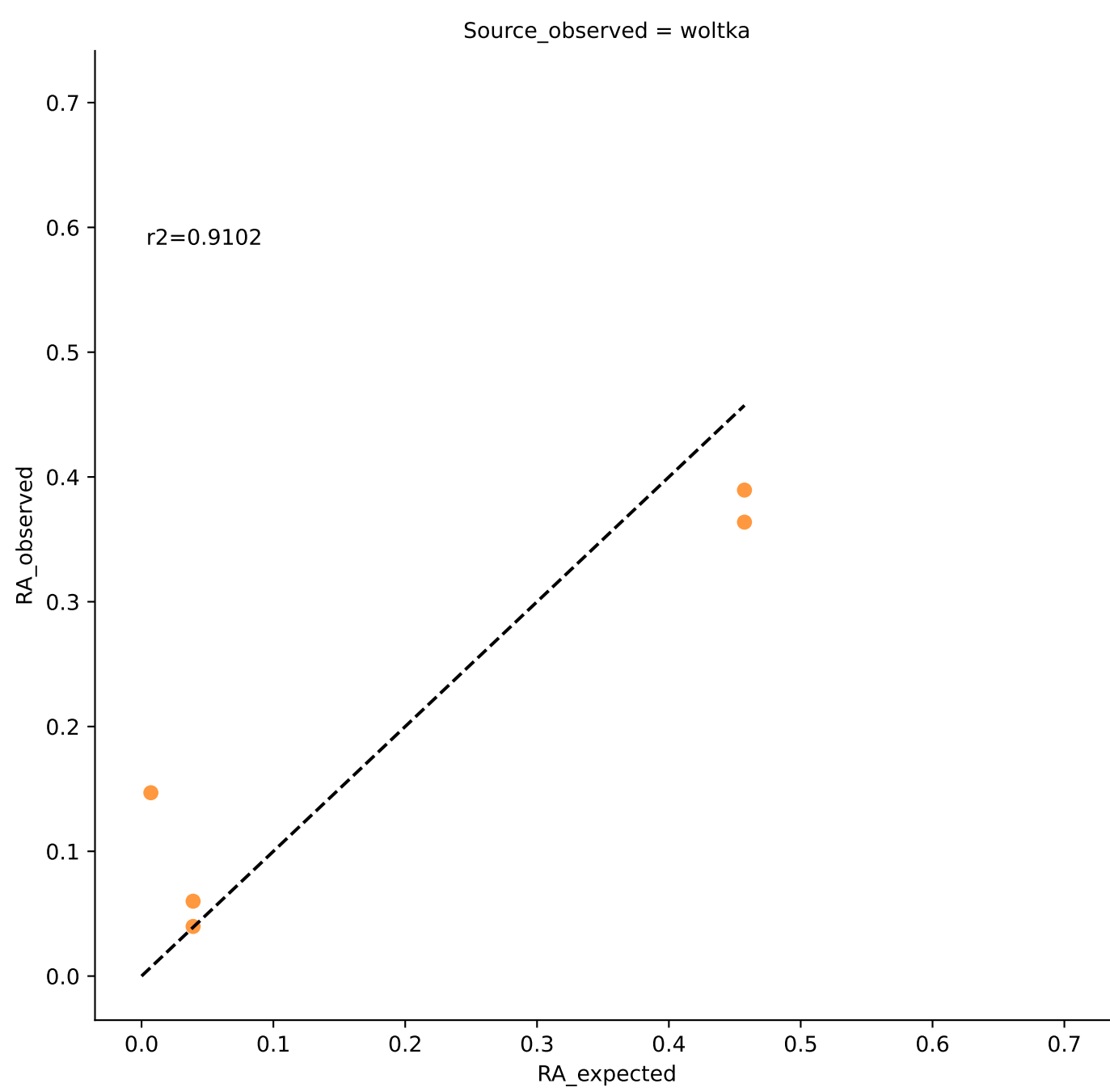
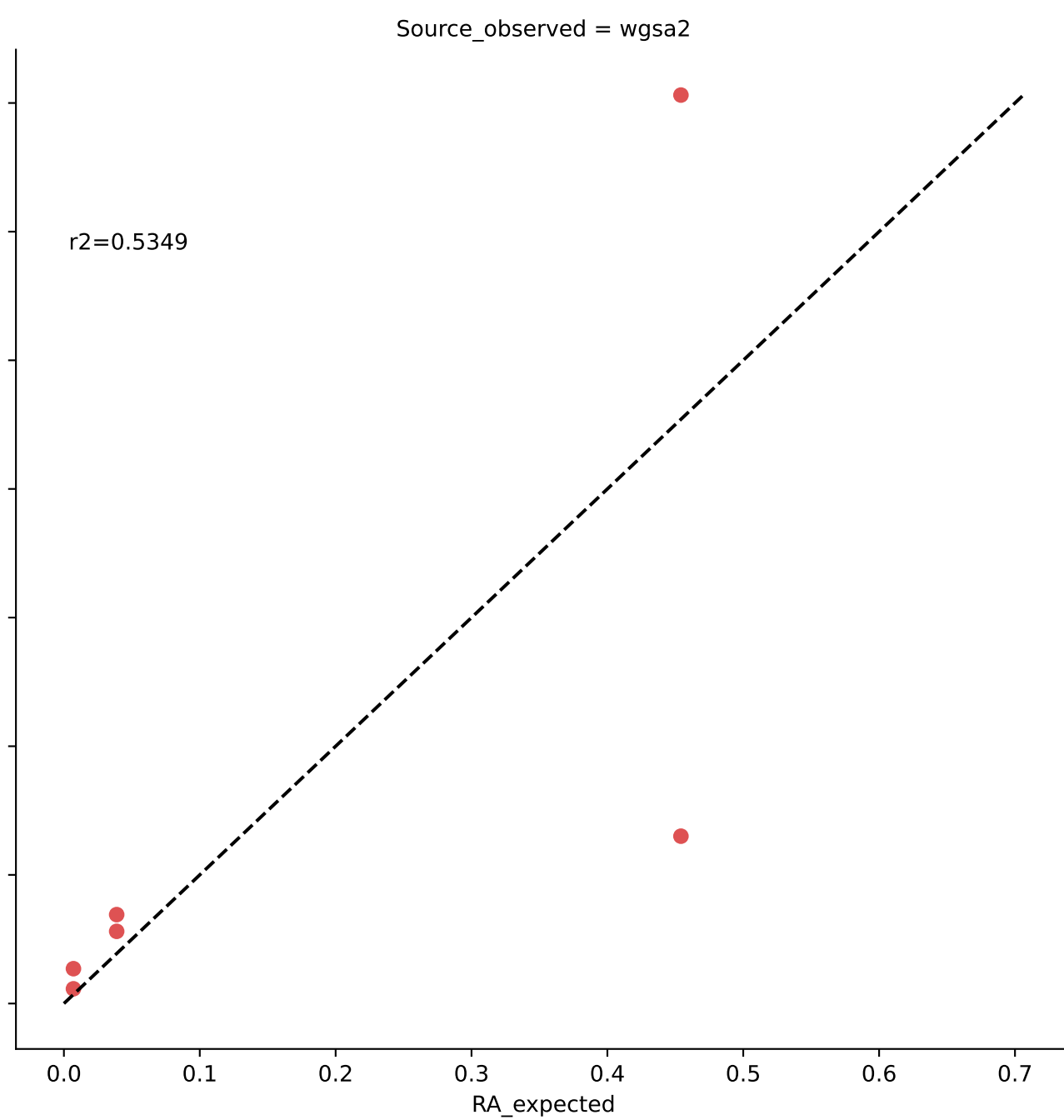
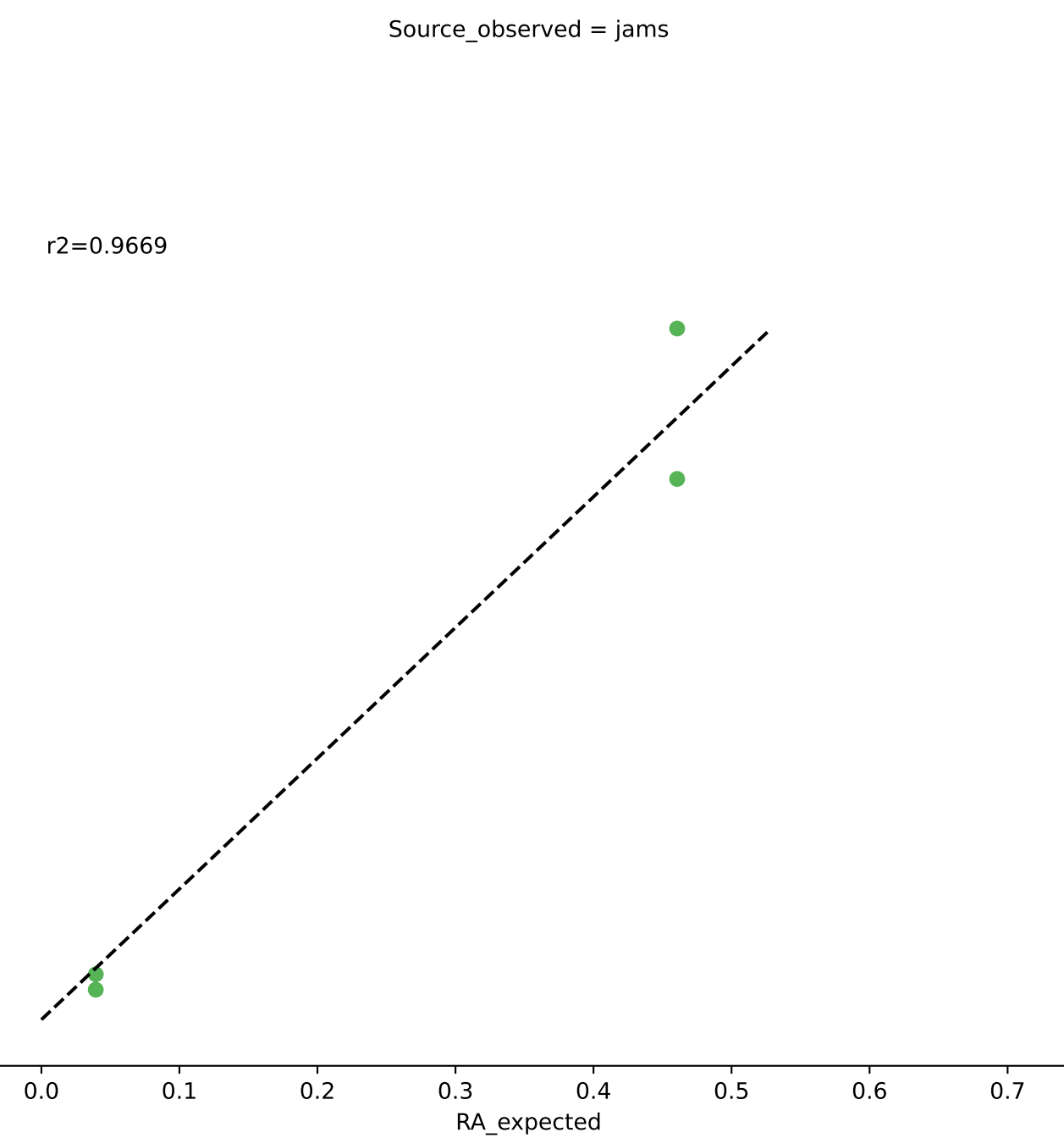
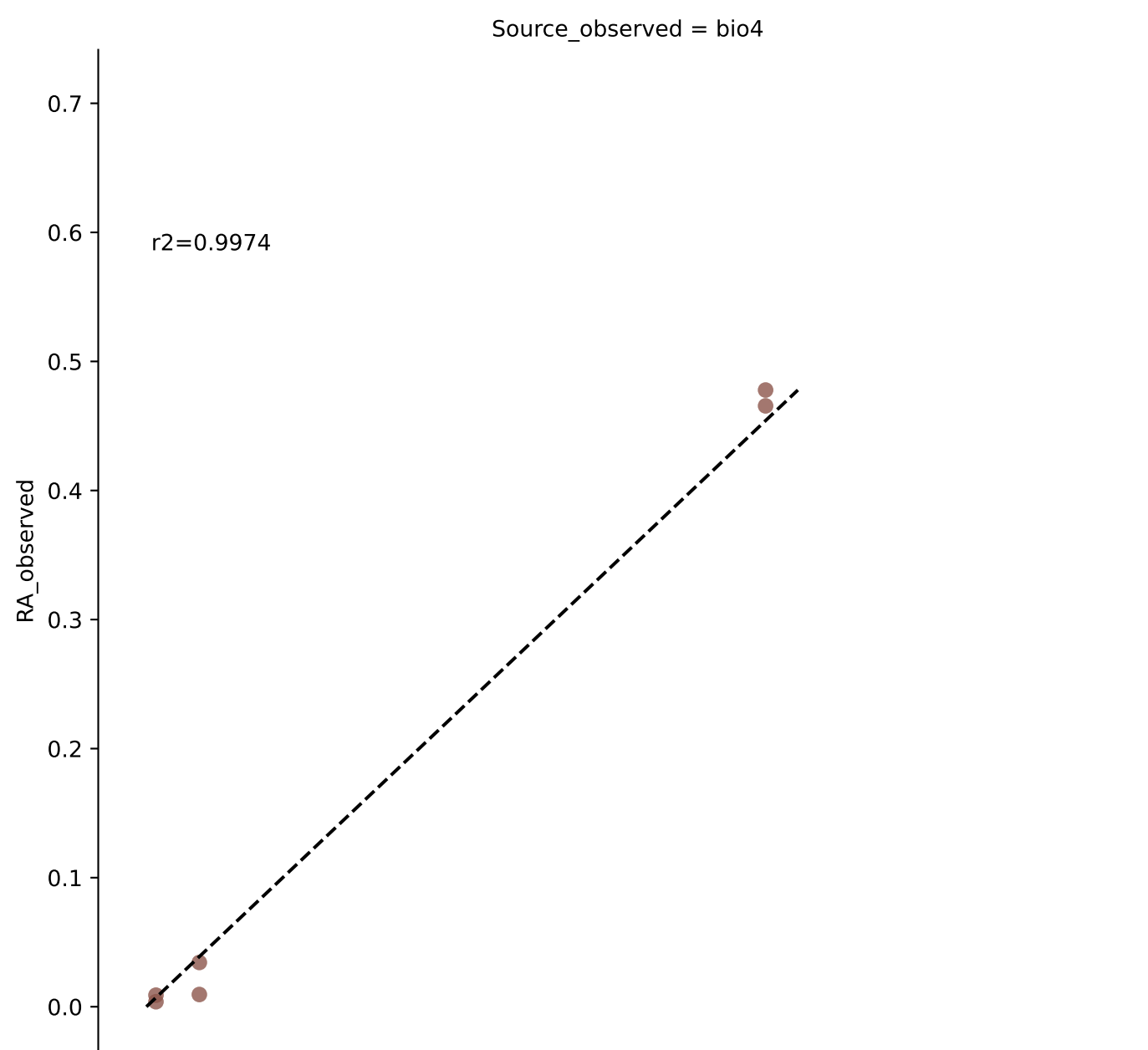


Bivariate Linear Regression for Sample EG in Experiment nist (Species at filter threshold 0.001)

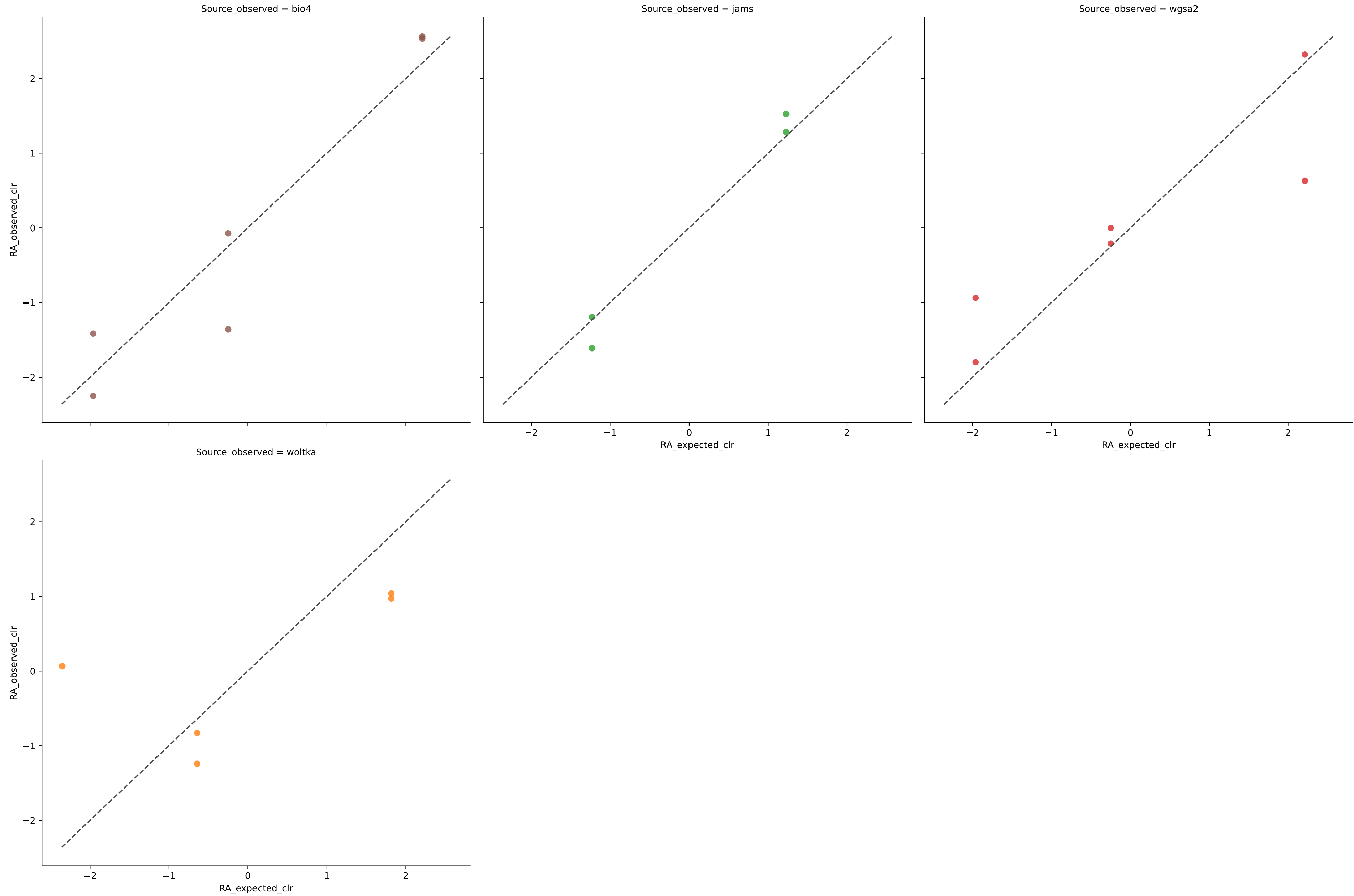


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	10	0.5696	0.0268	1.7058	0.8661	0.0342	100.0000	0.0000
jams	10	0.5145	0.0272	3.2110	0.8638	0.0390	100.0000	0.0000
wgsa2	10	0.3169	0.0460	2.3094	0.7702	0.0568	100.0000	0.0000
woltka	9	0.2616	0.0812	3.1357	0.6347	0.0973	100.0000	0.0000

Bivariate Linear Regression for Sample MIX-A in Experiment nist (Species at filter threshold 0.001)

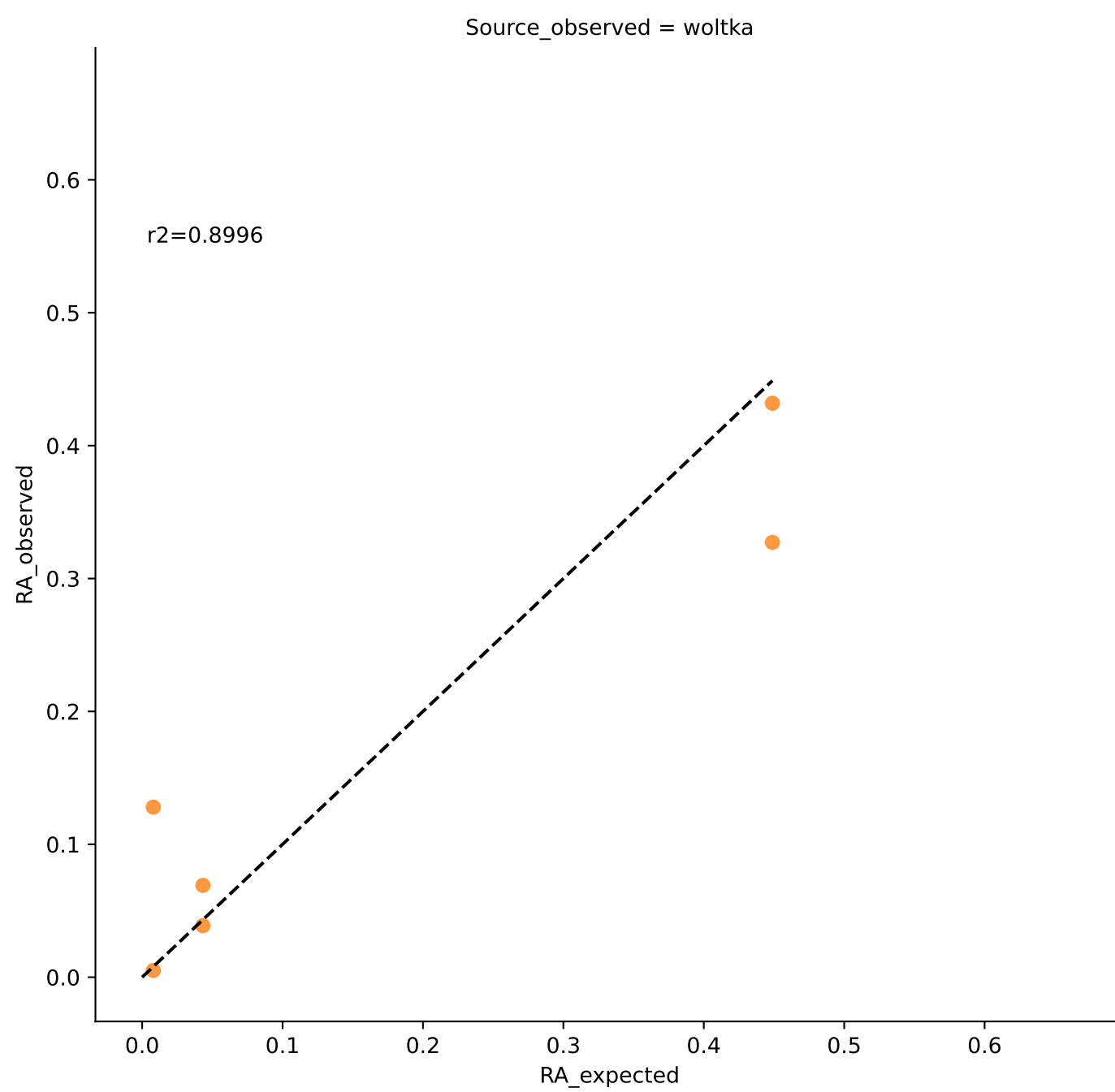
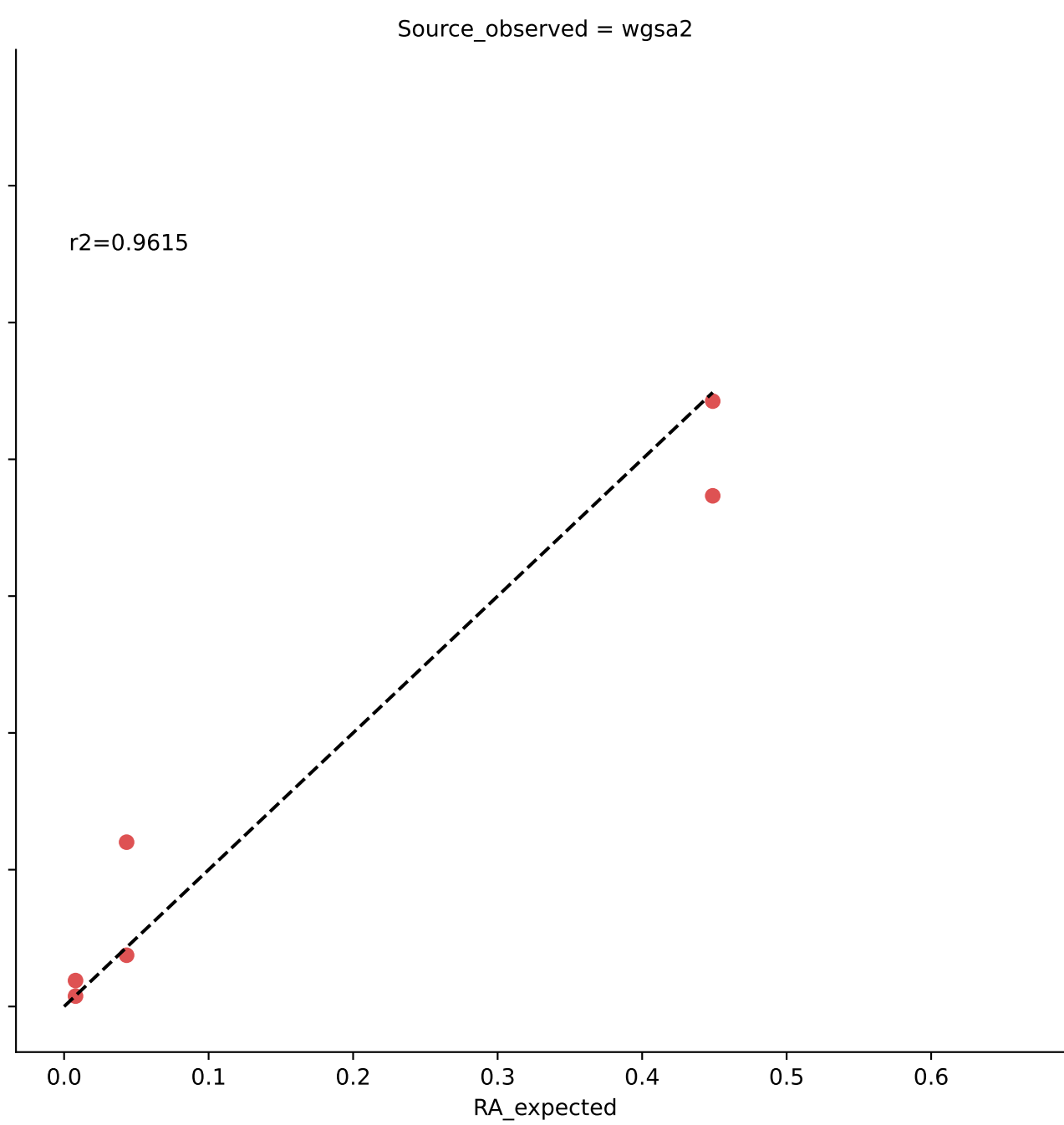
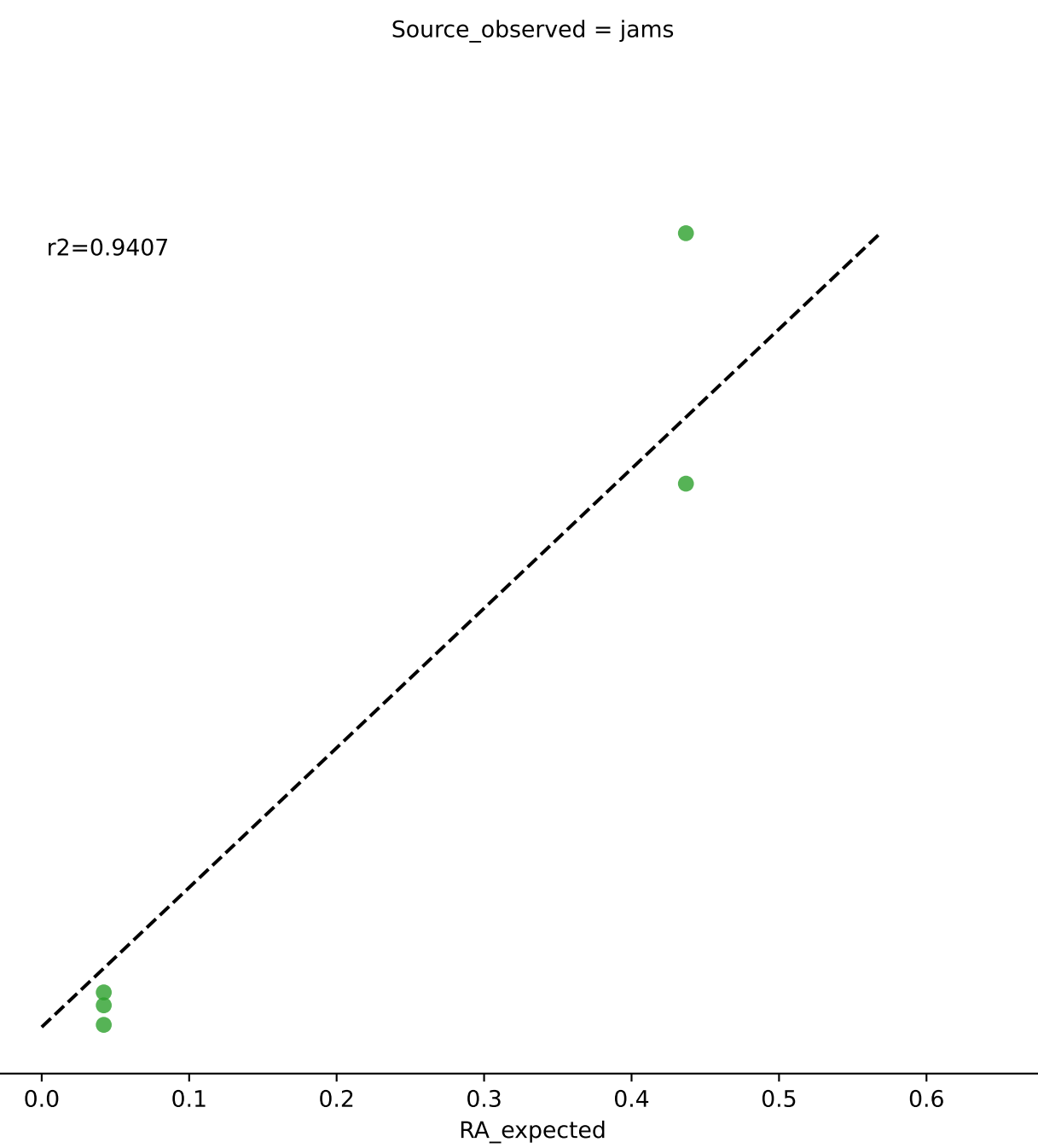
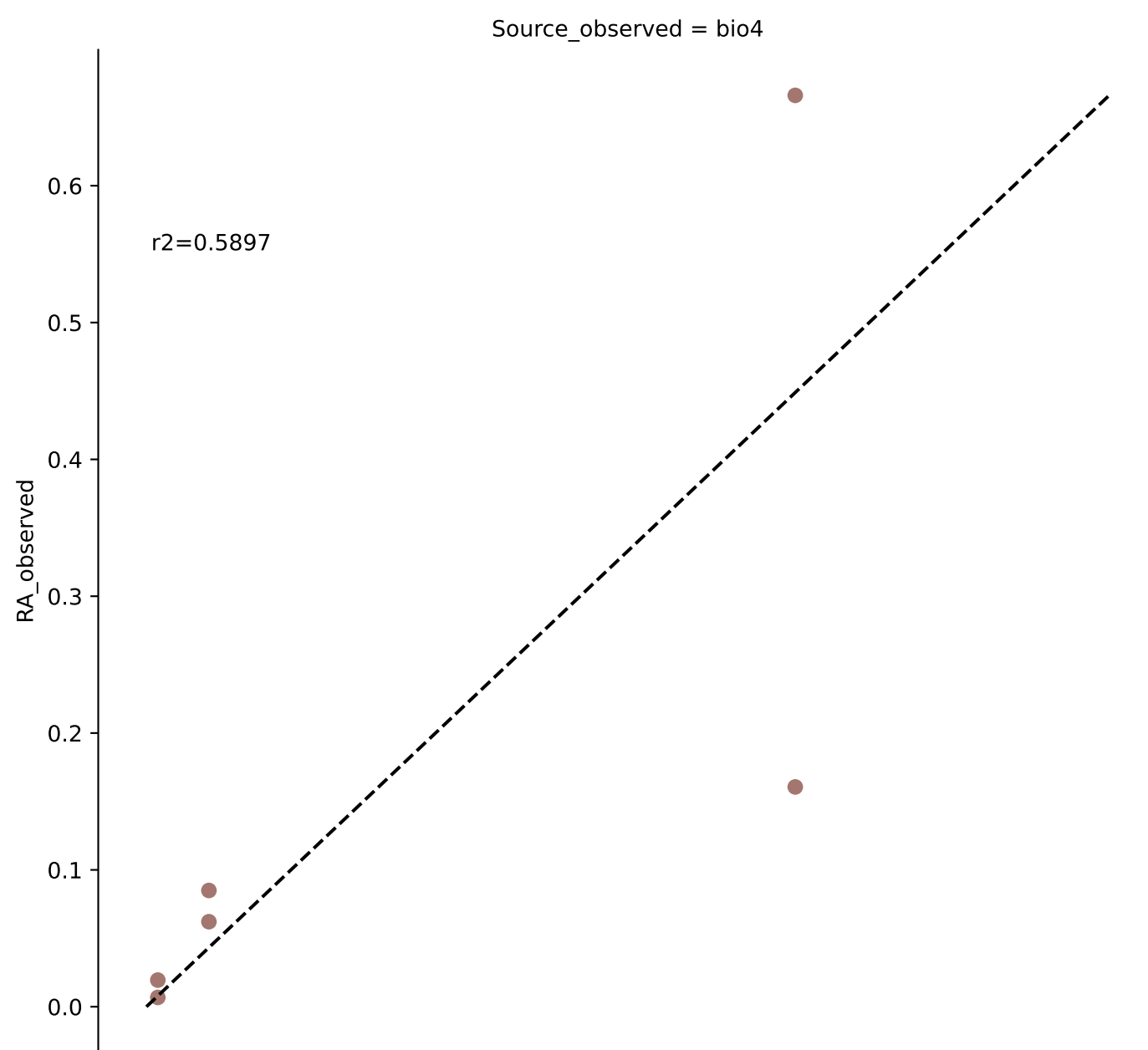


Bivariate Linear Regression for Sample MIX-A in Experiment nist (Species at filter threshold 0.001)

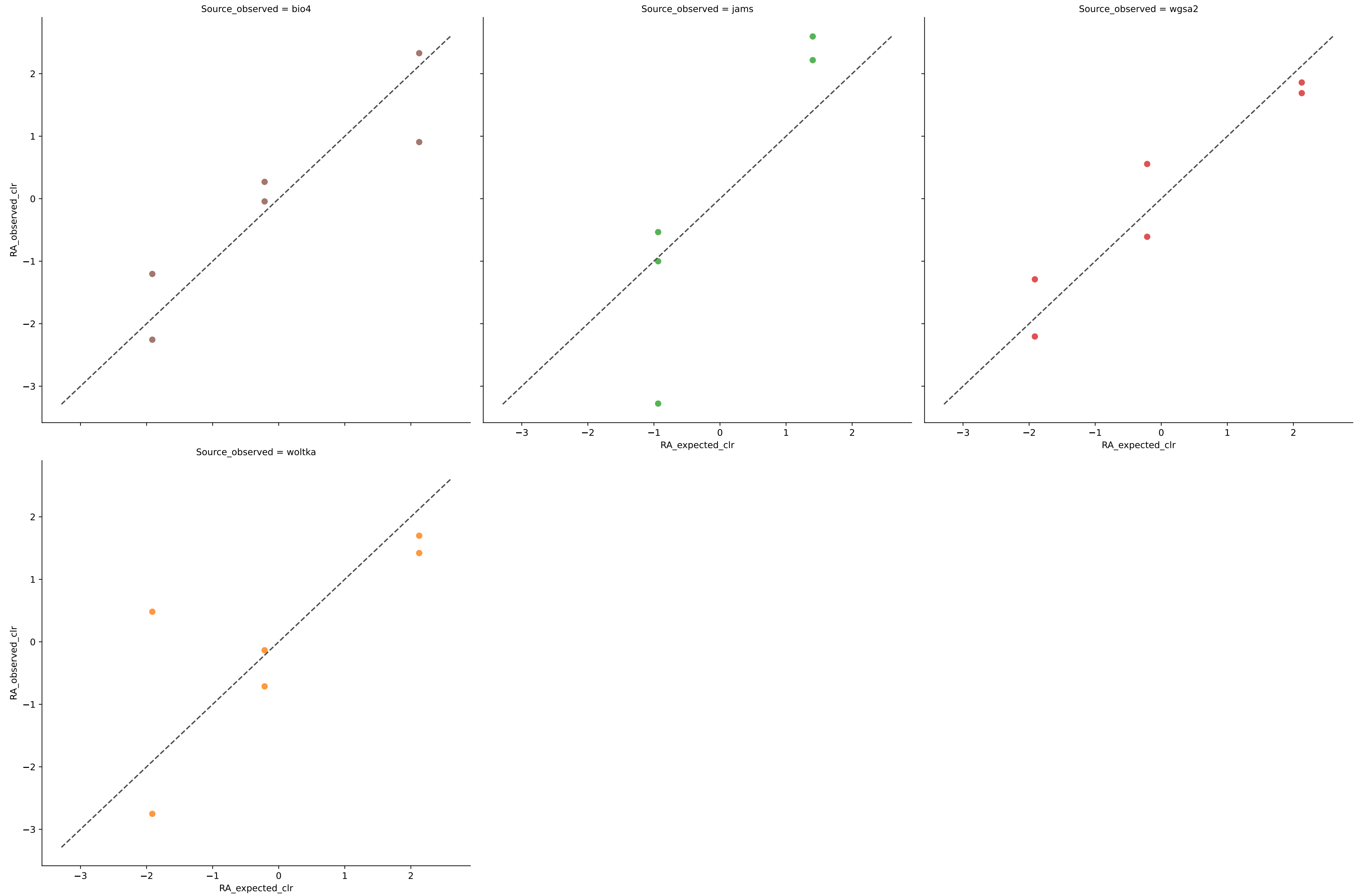


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	6	0.9974	0.0124	1.3695	0.9629	0.0163	100.0000	0.0000
jams	4	0.9669	0.0341	0.4885	0.9319	0.0422	100.0000	0.0000
wgsa2	6	0.5349	0.1080	1.9072	0.6759	0.1684	100.0000	0.0000
woltka	5	0.9102	0.0646	2.7491	0.8386	0.0817	100.0000	0.0000

Bivariate Linear Regression for Sample MIX-B in Experiment nist (Species at filter threshold 0.001)

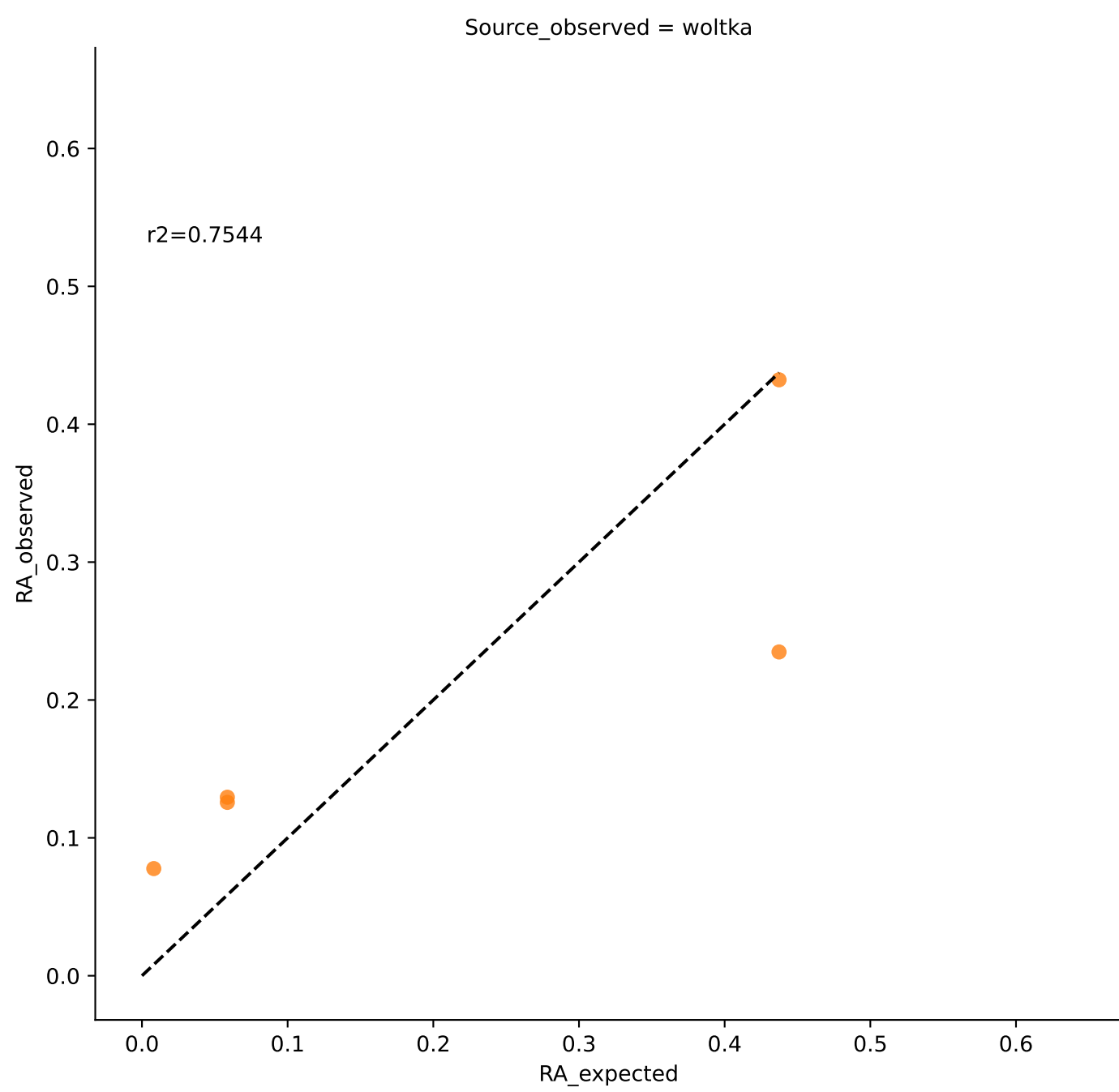
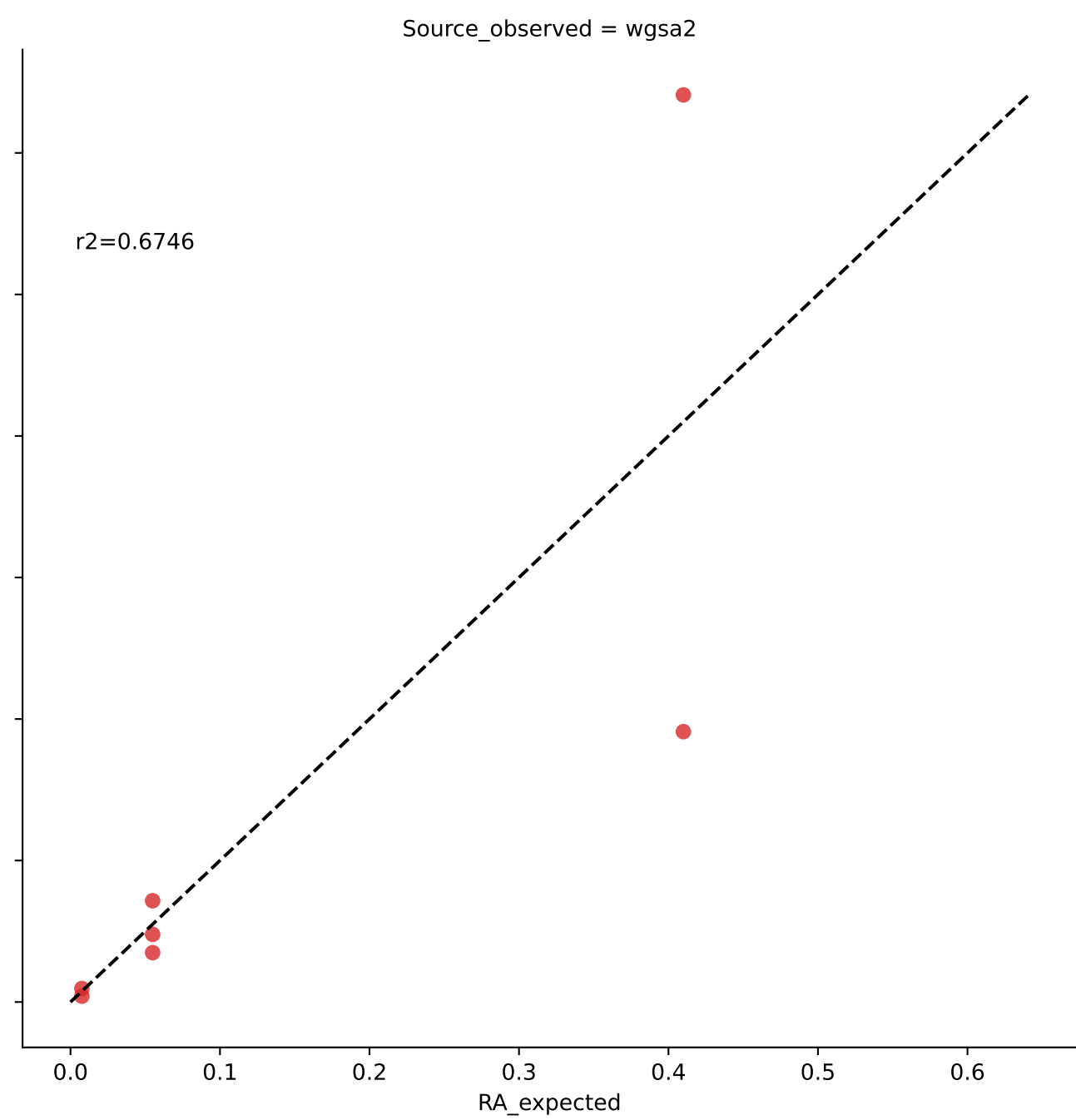
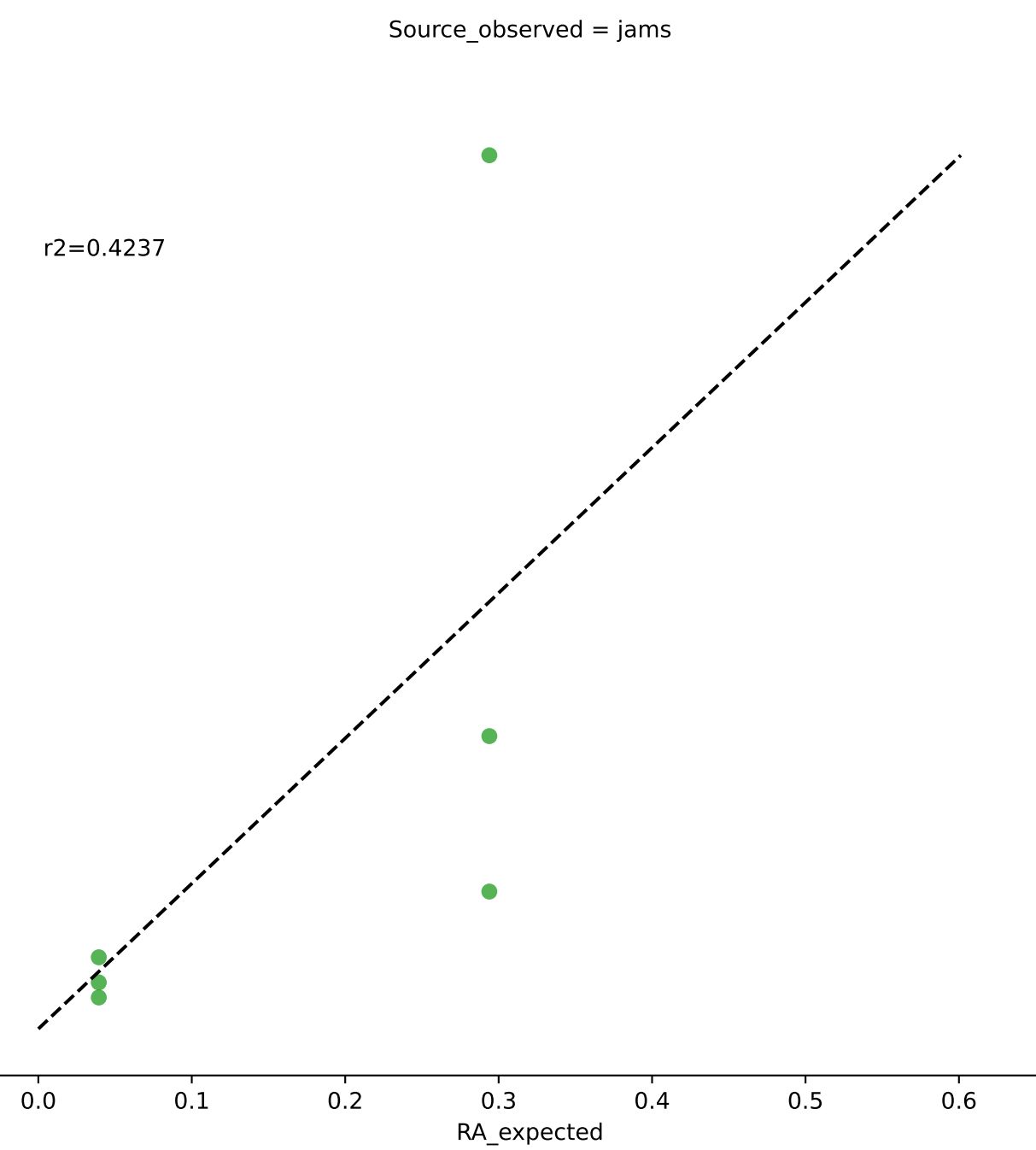
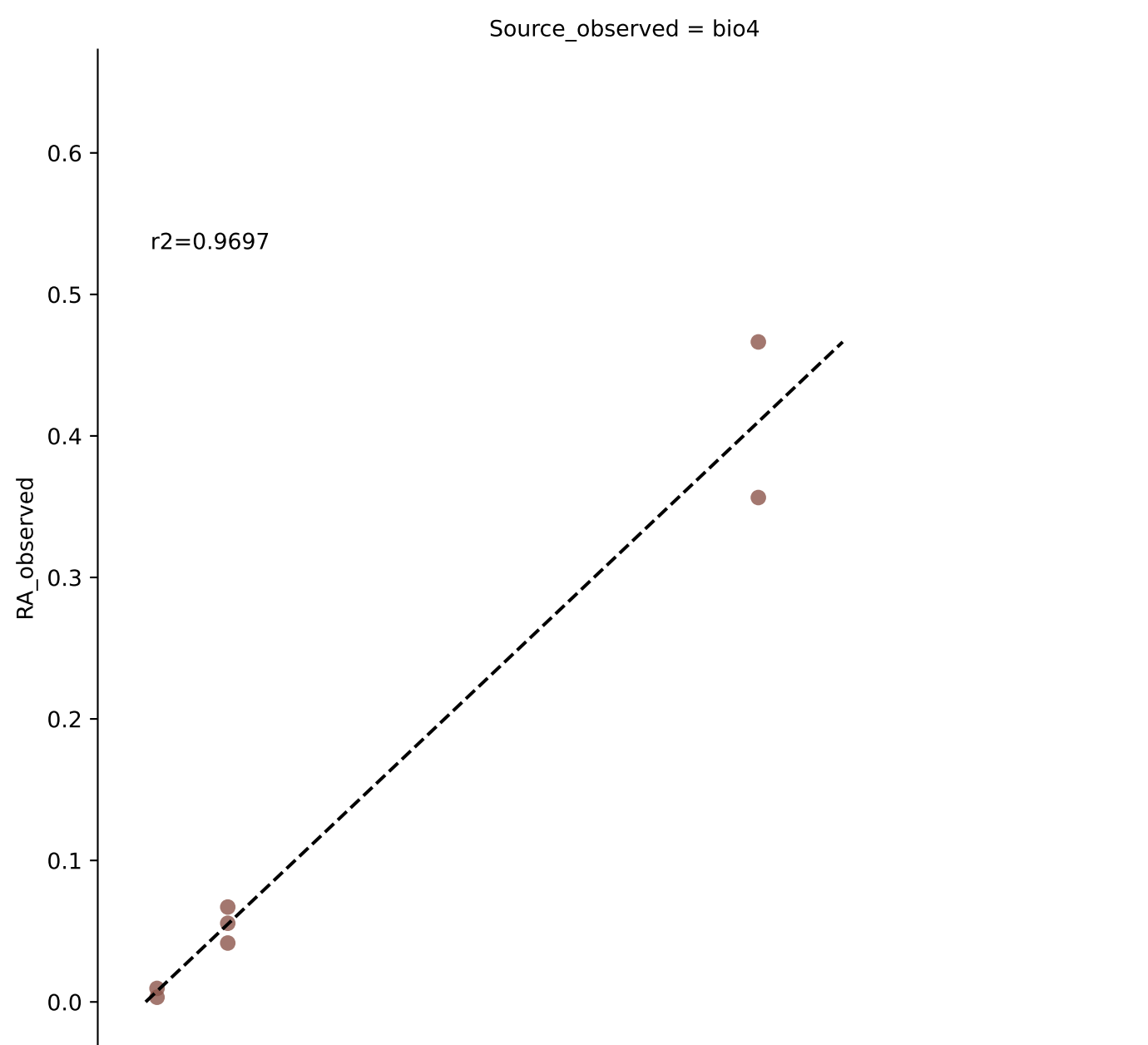


Bivariate Linear Regression for Sample MIX-B in Experiment nist (Species at filter threshold 0.001)

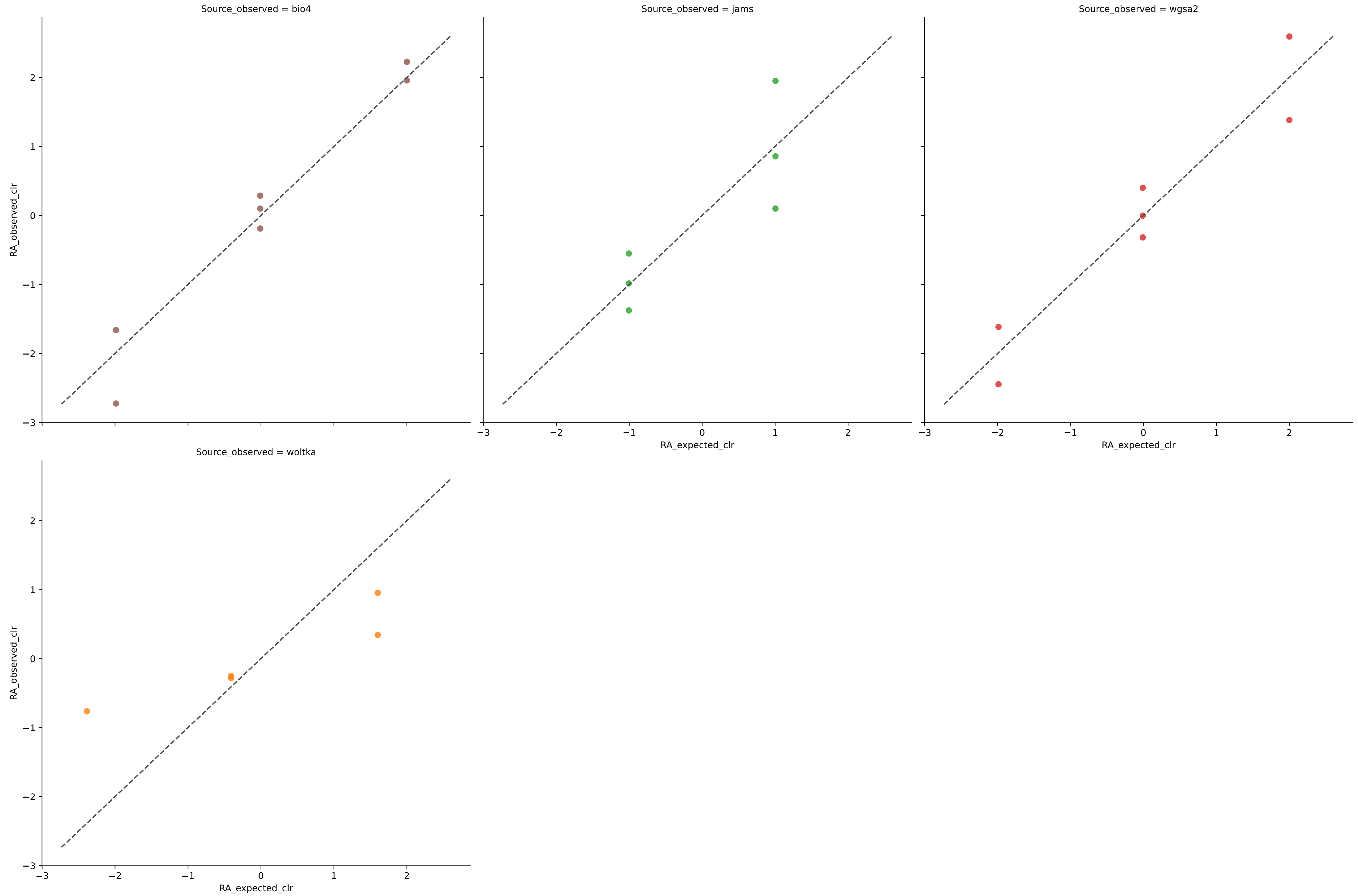


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	6	0.5897	0.0964	1.5530	0.7107	0.1486	100.0000	0.0000
jams	5	0.9407	0.0527	2.7810	0.8682	0.0667	100.0000	0.0000
wgsa2	6	0.9615	0.0293	1.2180	0.9120	0.0444	100.0000	0.0000
woltka	6	0.8996	0.0486	2.7159	0.8541	0.0709	100.0000	0.0000

Bivariate Linear Regression for Sample MIX-C in Experiment nist (Species at filter threshold 0.001)

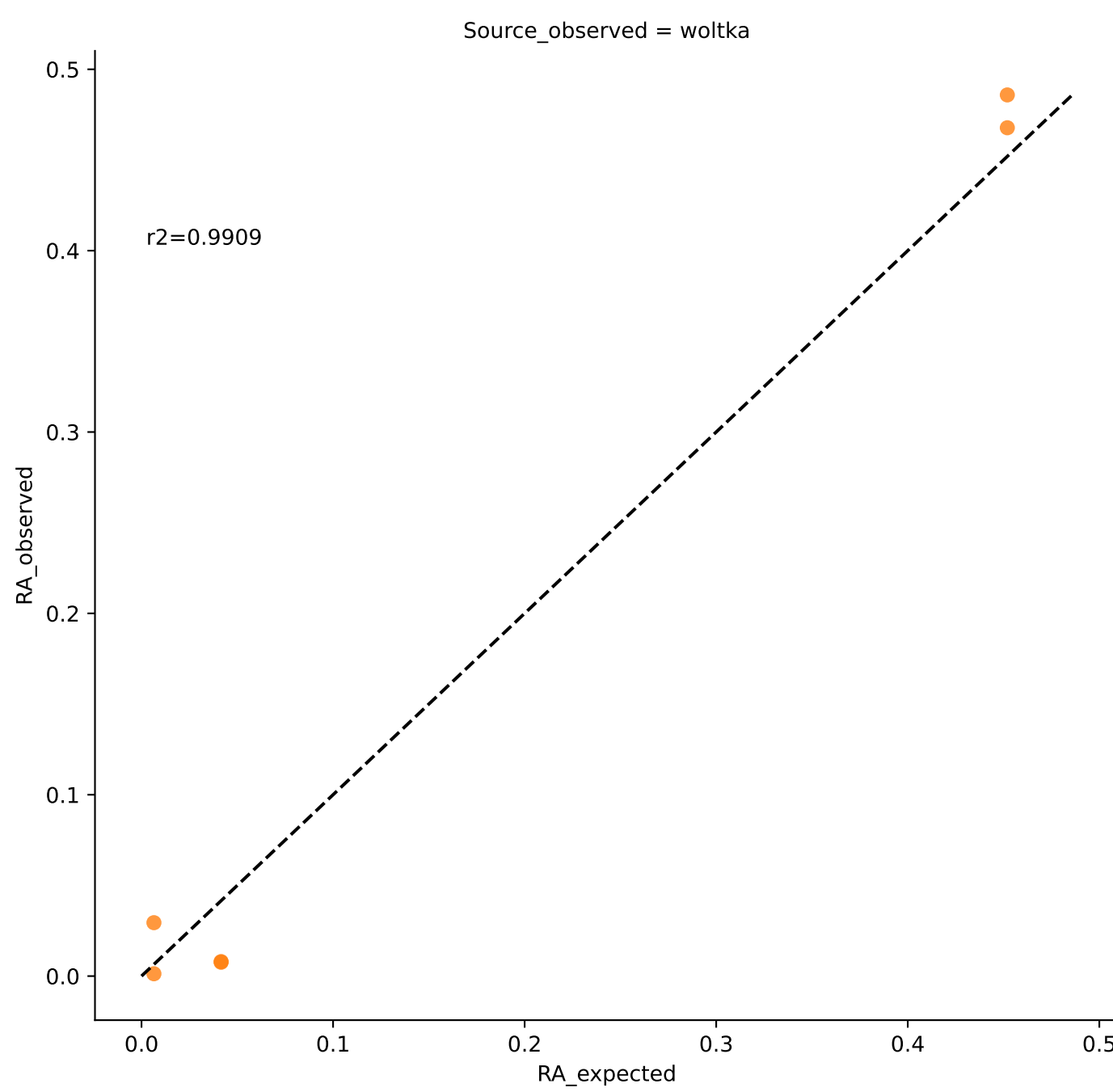
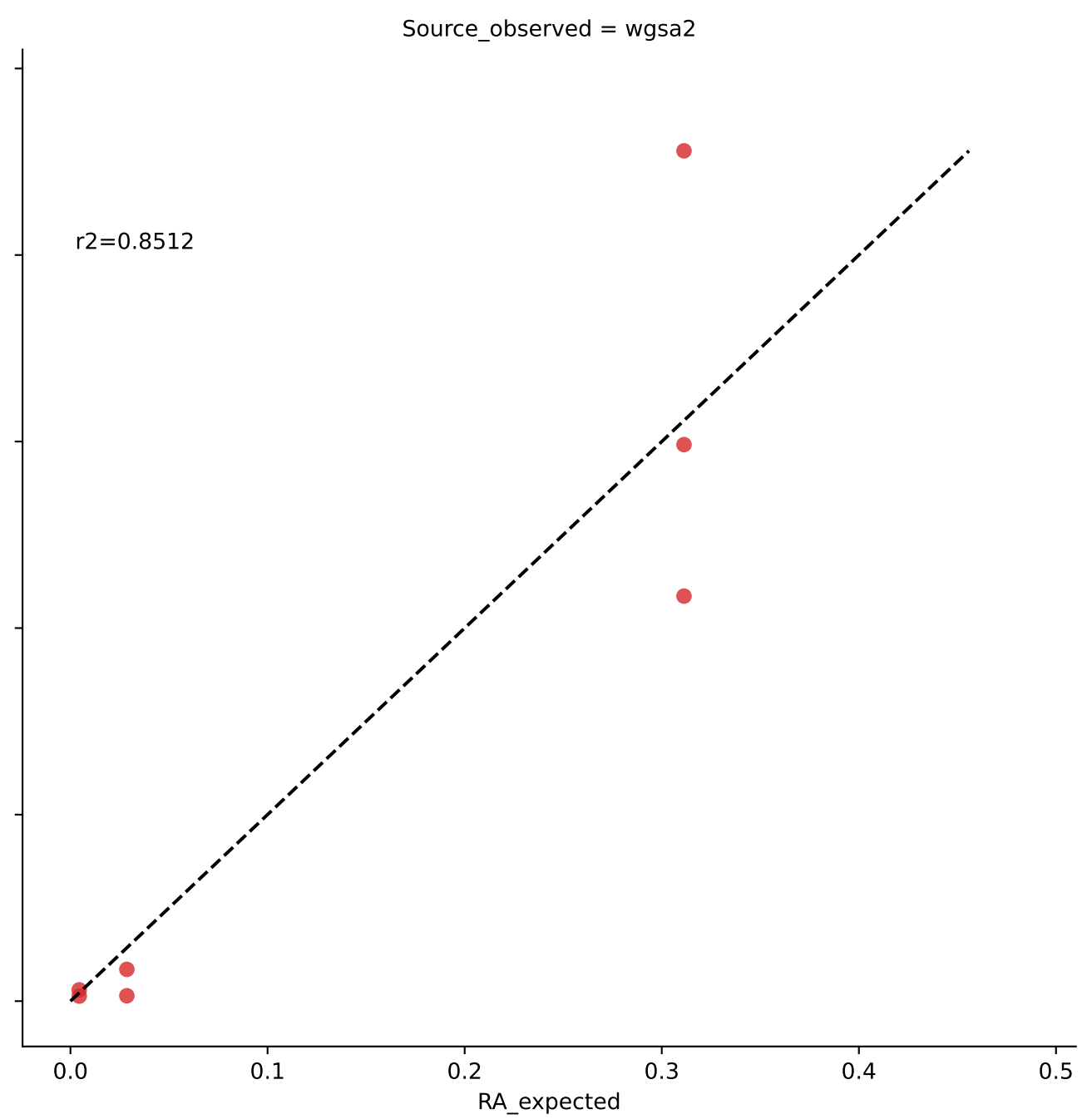
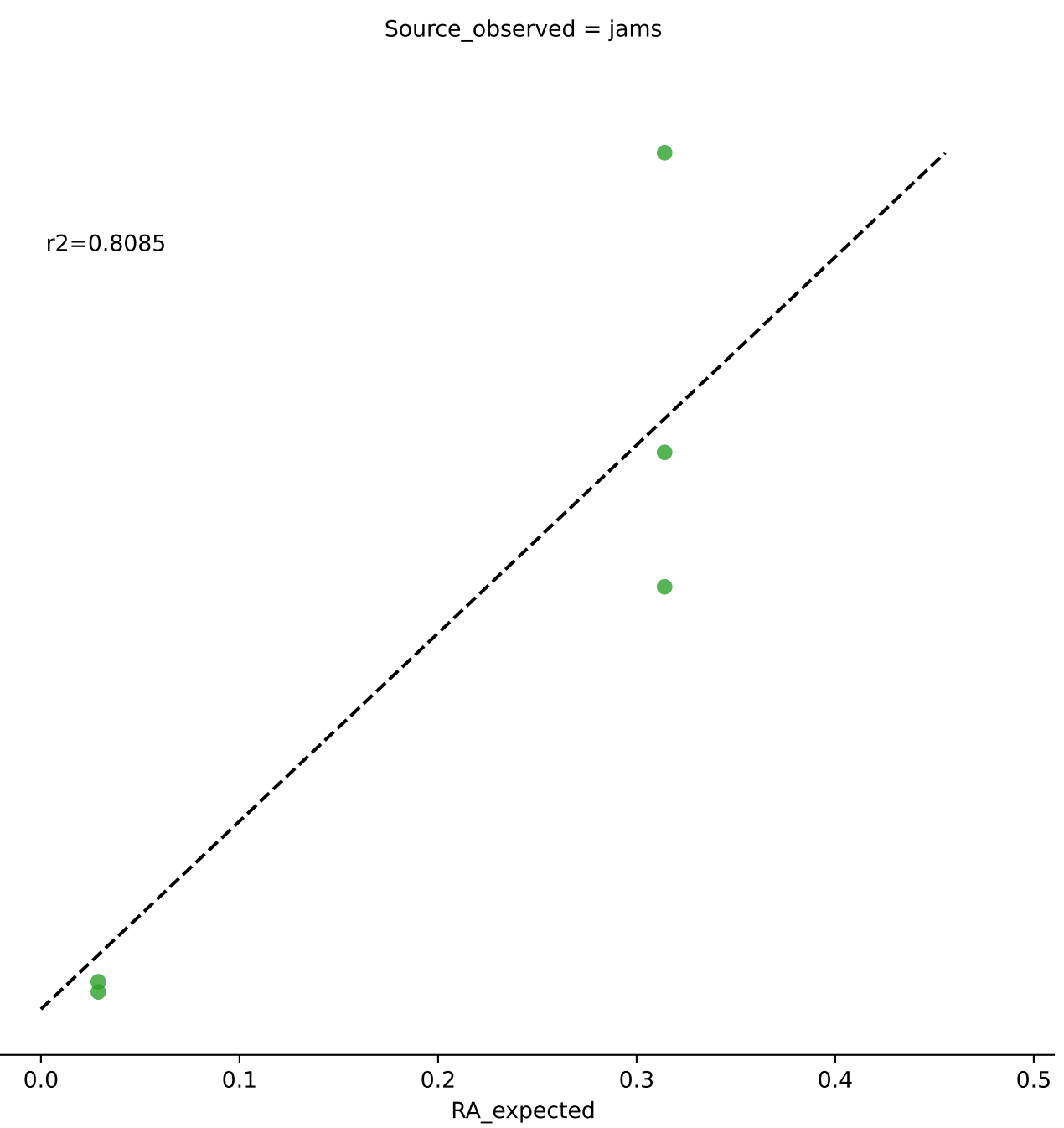
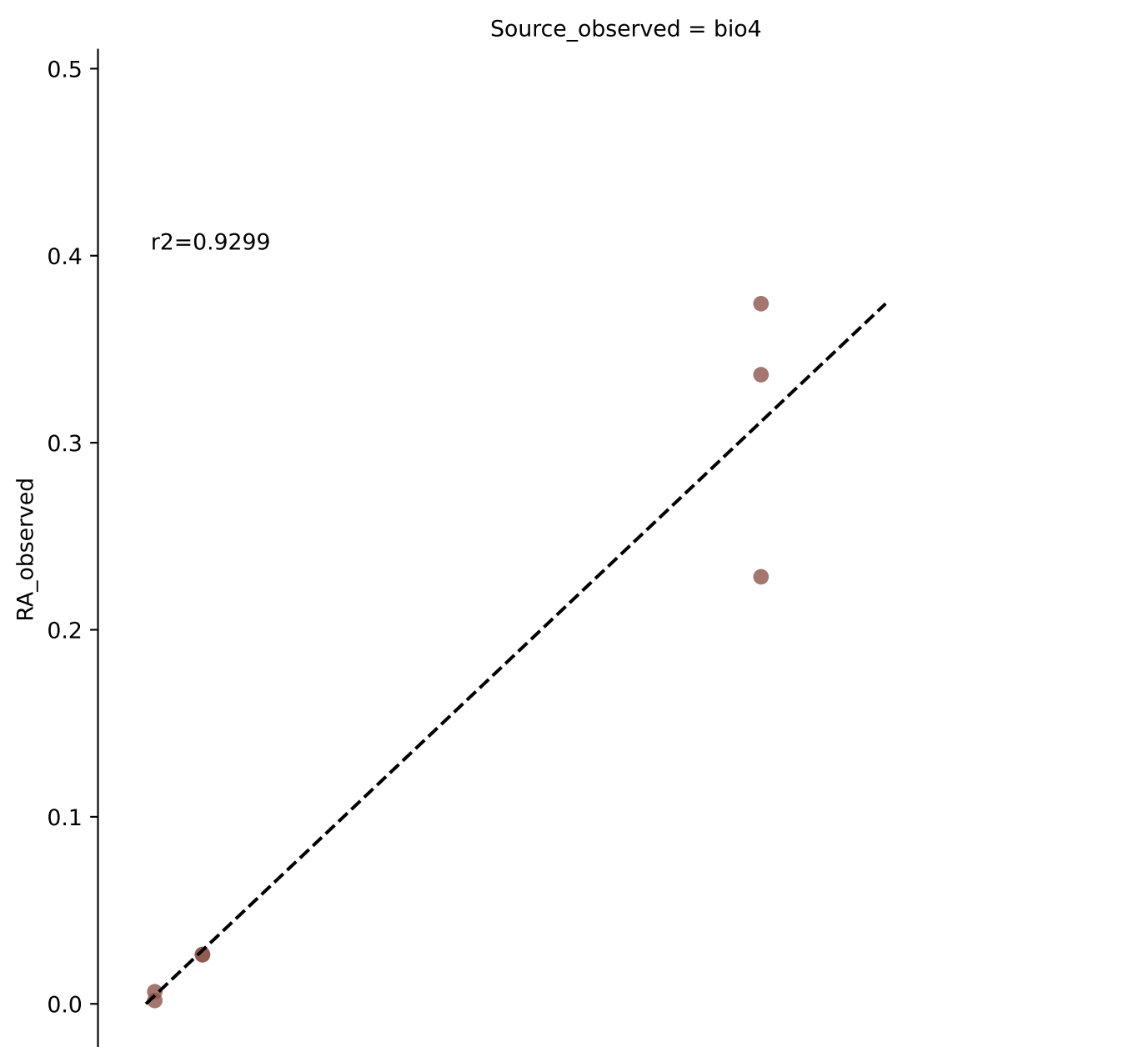


Bivariate Linear Regression for Sample MIX-C in Experiment nist (Species at filter threshold 0.001)

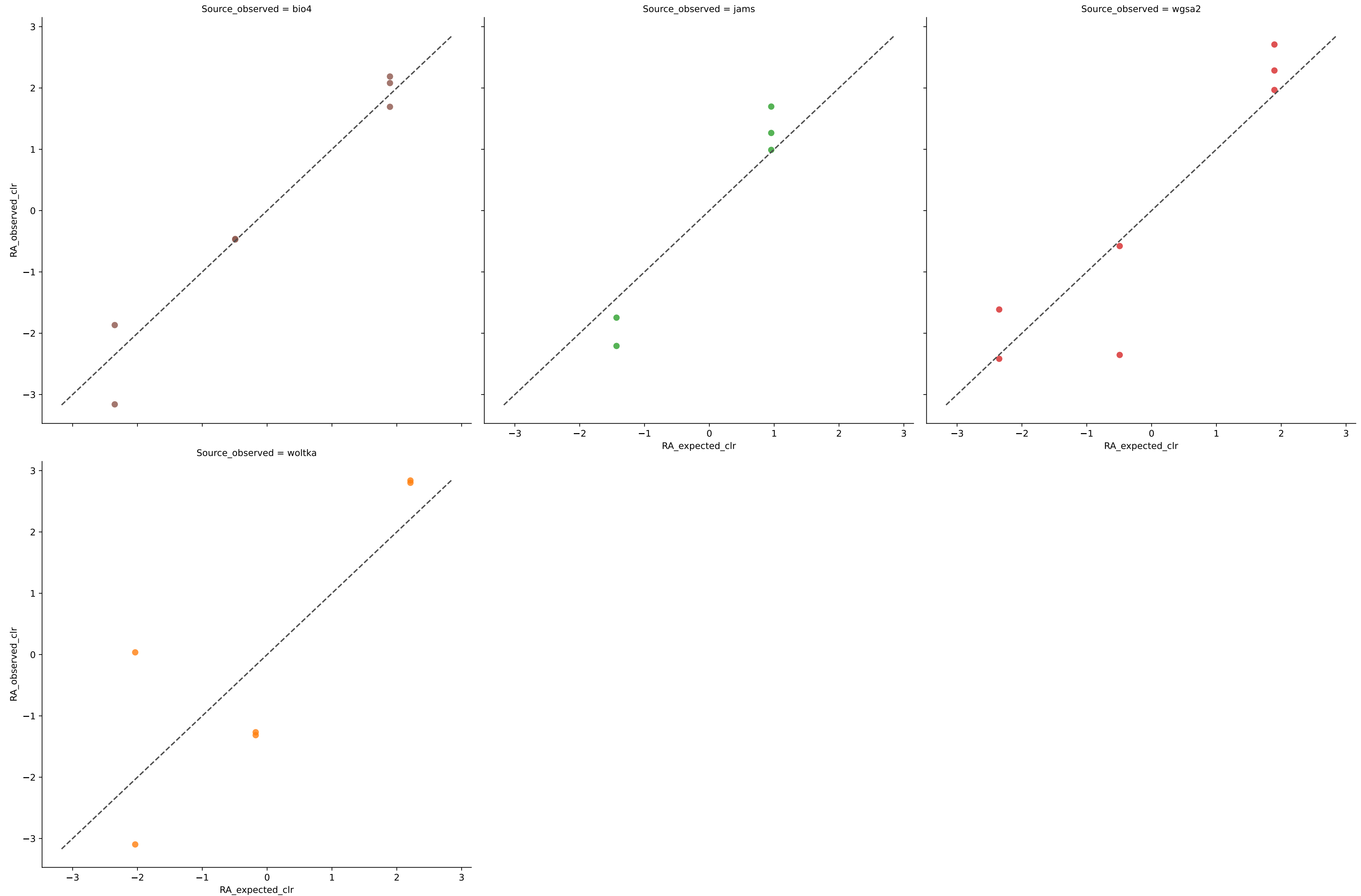


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	7	0.9697	0.0203	0.9148	0.9288	0.0302	100.0000	0.0000
jams	6	0.4237	0.1057	1.4416	0.6829	0.1545	100.0000	0.0000
wgsa2	7	0.6746	0.0713	1.1599	0.7504	0.1207	100.0000	0.0000
woltka	5	0.7544	0.0830	2.1628	0.7924	0.1053	100.0000	0.0000

Bivariate Linear Regression for Sample MIX-D in Experiment nist (Species at filter threshold 0.001)

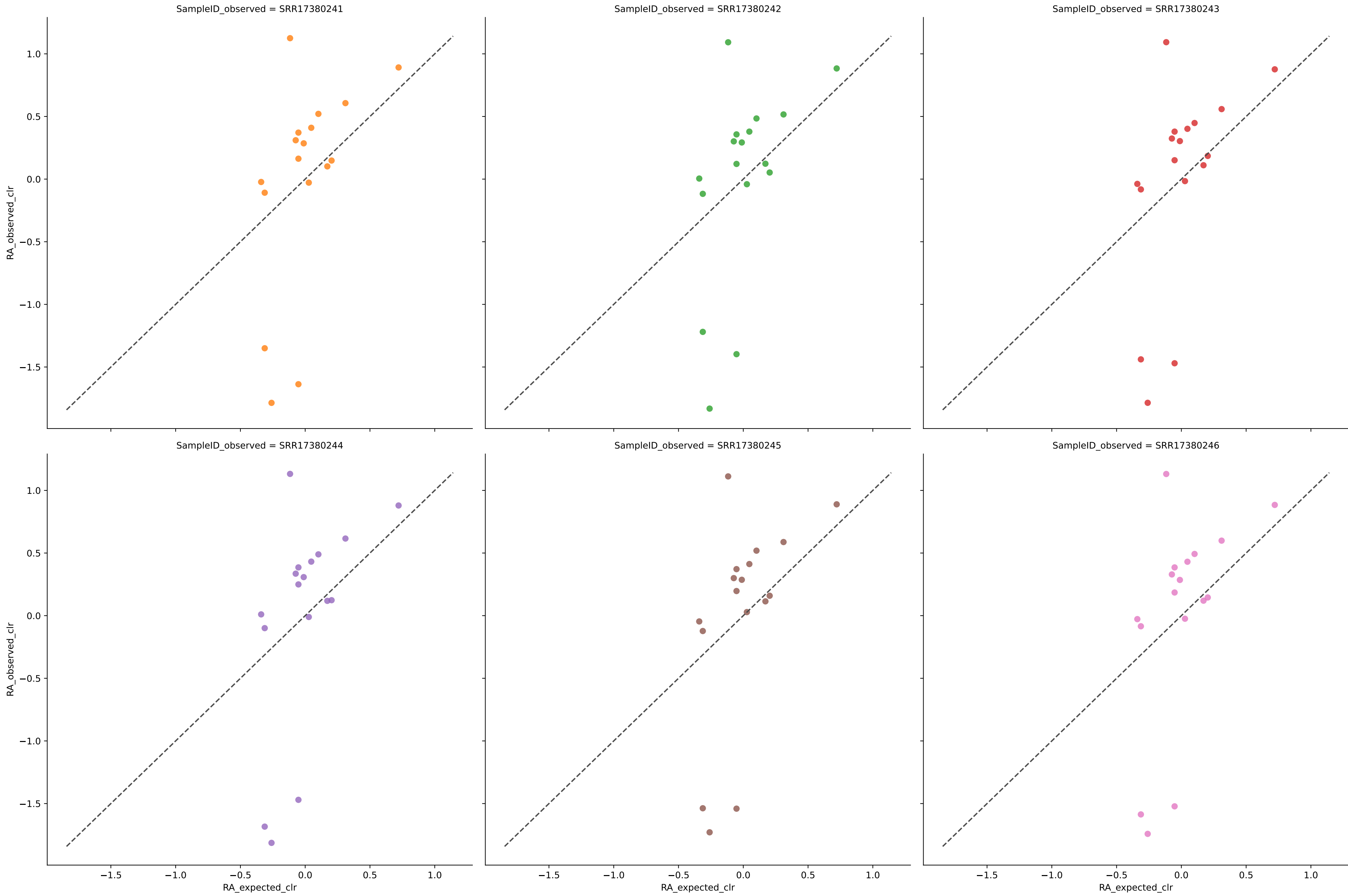


Bivariate Linear Regression for Sample MIX-D in Experiment nist (Species at filter threshold 0.001)



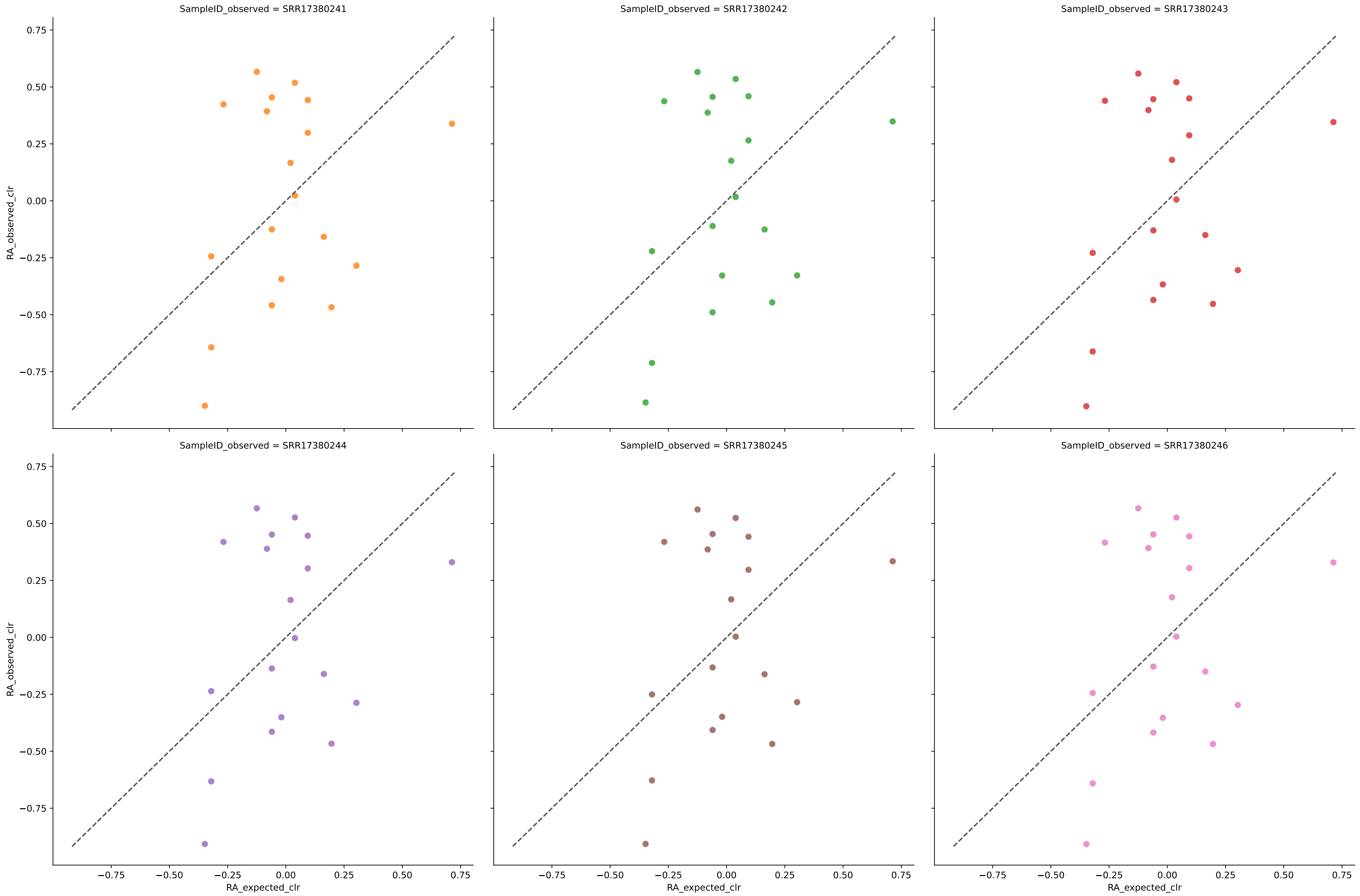
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	7	0.9299	0.0258	1.0255	0.9098	0.0406	100.0000	0.0000
jams	5	0.8085	0.0566	1.1619	0.8586	0.0760	100.0000	0.0000
wgsa2	7	0.8512	0.0418	2.2012	0.8538	0.0663	100.0000	0.0000
woltka	6	0.9909	0.0242	2.9407	0.9273	0.0266	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using bio4 in Experiment tourlousse with filter 0.001



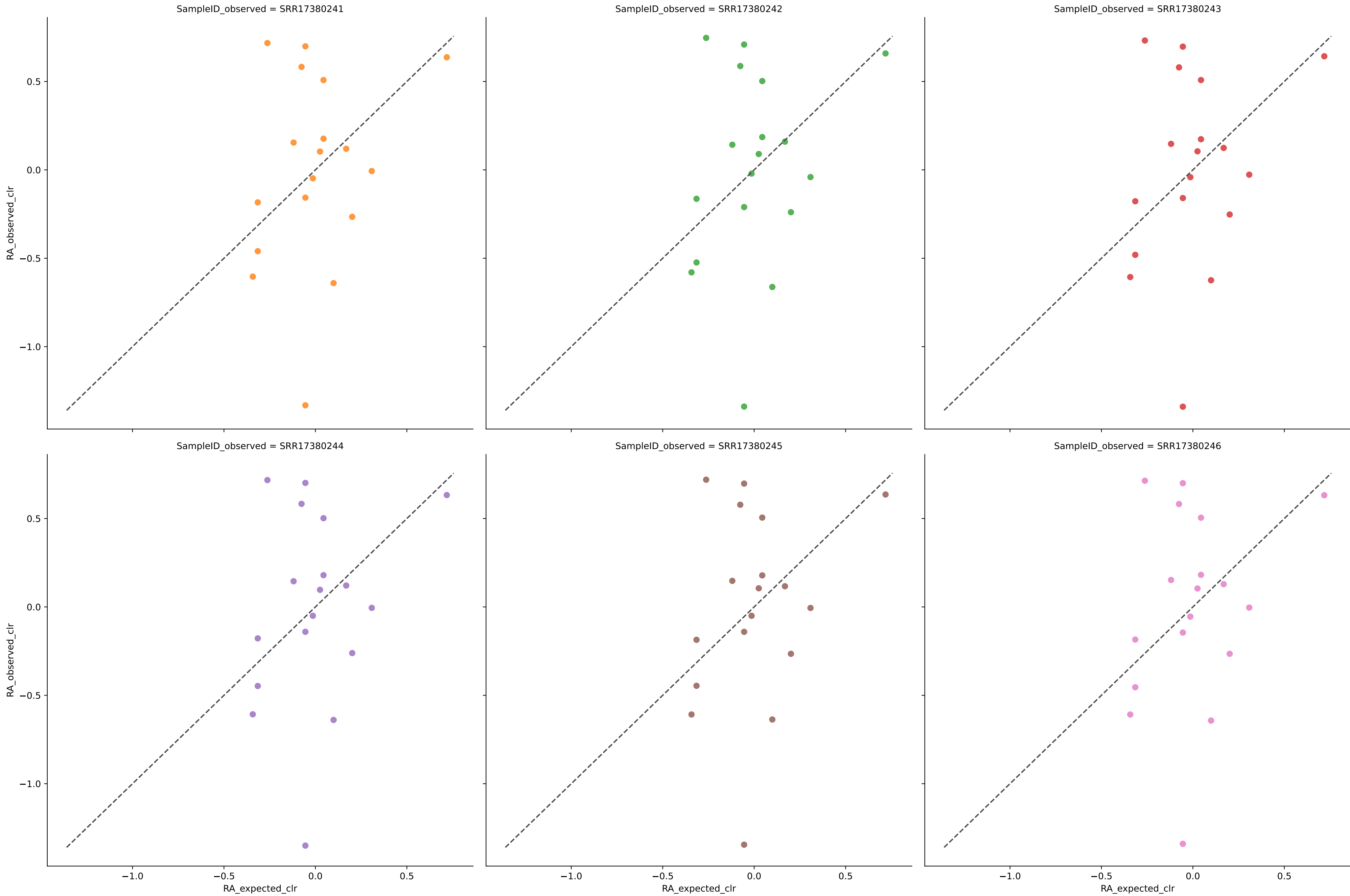
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR17380241	17	0.2683	0.0031	2.9154	0.8404	0.0048	100.0000	0.0000
SRR17380242	17	0.2667	0.0030	2.7401	0.8446	0.0047	100.0000	0.0000
SRR17380243	17	0.2744	0.0030	2.8350	0.8461	0.0046	100.0000	0.0000
SRR17380244	17	0.2551	0.0032	3.0019	0.8349	0.0048	100.0000	0.0000
SRR17380245	17	0.2771	0.0031	2.8973	0.8429	0.0047	100.0000	0.0000
SRR17380246	17	0.2608	0.0032	2.9331	0.8386	0.0048	100.0000	0.0000
Average	17	0.2671	0.0031	2.8871	0.8412	0.0048	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using jams in Experiment tourlousse with filter 0.001



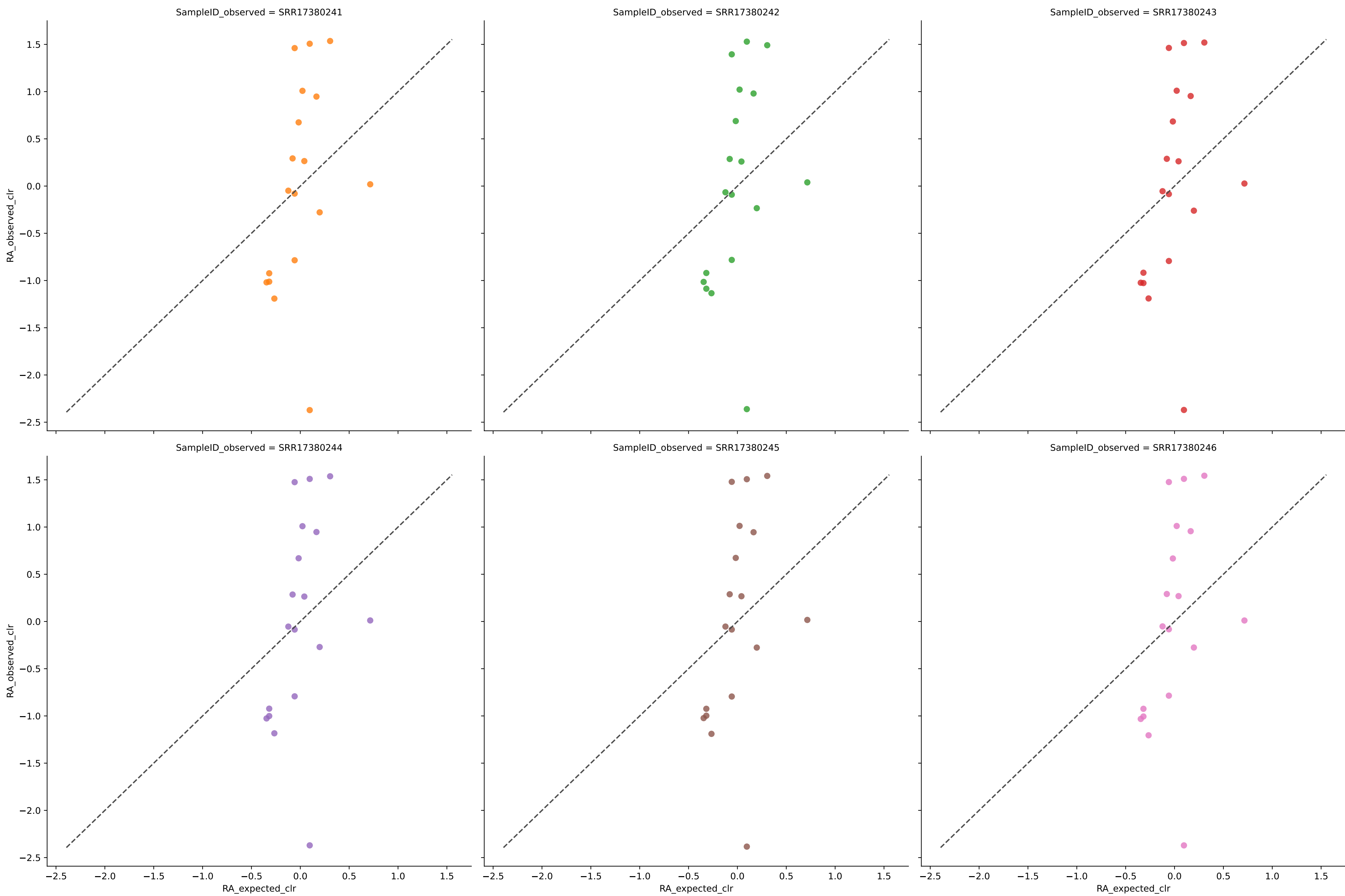
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR17380241	19	0.0294	0.0033	1.8893	0.8104	0.0039	100.0000	0.0000
SRR17380242	19	0.0298	0.0033	1.9110	0.8093	0.0039	100.0000	0.0000
SRR17380243	19	0.0296	0.0033	1.8964	0.8091	0.0039	100.0000	0.0000
SRR17380244	19	0.0276	0.0033	1.8859	0.8093	0.0039	100.0000	0.0000
SRR17380245	19	0.0291	0.0033	1.8776	0.8108	0.0039	100.0000	0.0000
SRR17380246	19	0.0276	0.0033	1.8902	0.8094	0.0039	100.0000	0.0000
Average	19	0.0289	0.0033	1.8917	0.8097	0.0039	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using wgsa in Experiment tourlousse with filter 0.001



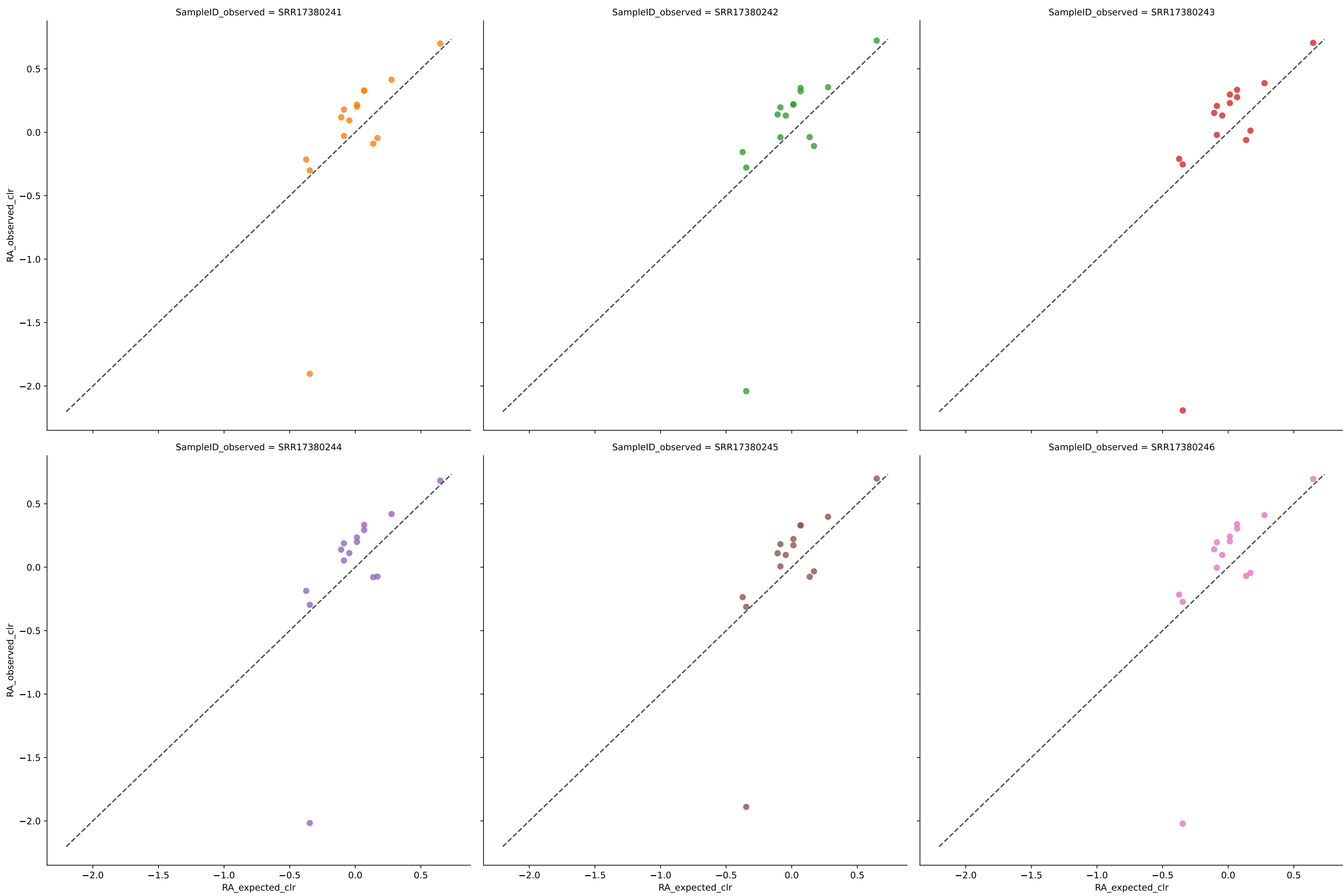
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR17380241	18	0.0730	0.0034	2.2129	0.8156	0.0044	100.0000	0.0000
SRR17380242	18	0.0747	0.0035	2.2448	0.8130	0.0045	100.0000	0.0000
SRR17380243	18	0.0729	0.0034	2.2182	0.8155	0.0044	100.0000	0.0000
SRR17380244	18	0.0714	0.0034	2.2201	0.8158	0.0044	100.0000	0.0000
SRR17380245	18	0.0727	0.0034	2.2160	0.8165	0.0044	100.0000	0.0000
SRR17380246	18	0.0723	0.0034	2.2157	0.8160	0.0044	100.0000	0.0000
Average	18	0.0728	0.0034	2.2213	0.8154	0.0044	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using woltka in Experiment tourlousse with filter 0.001



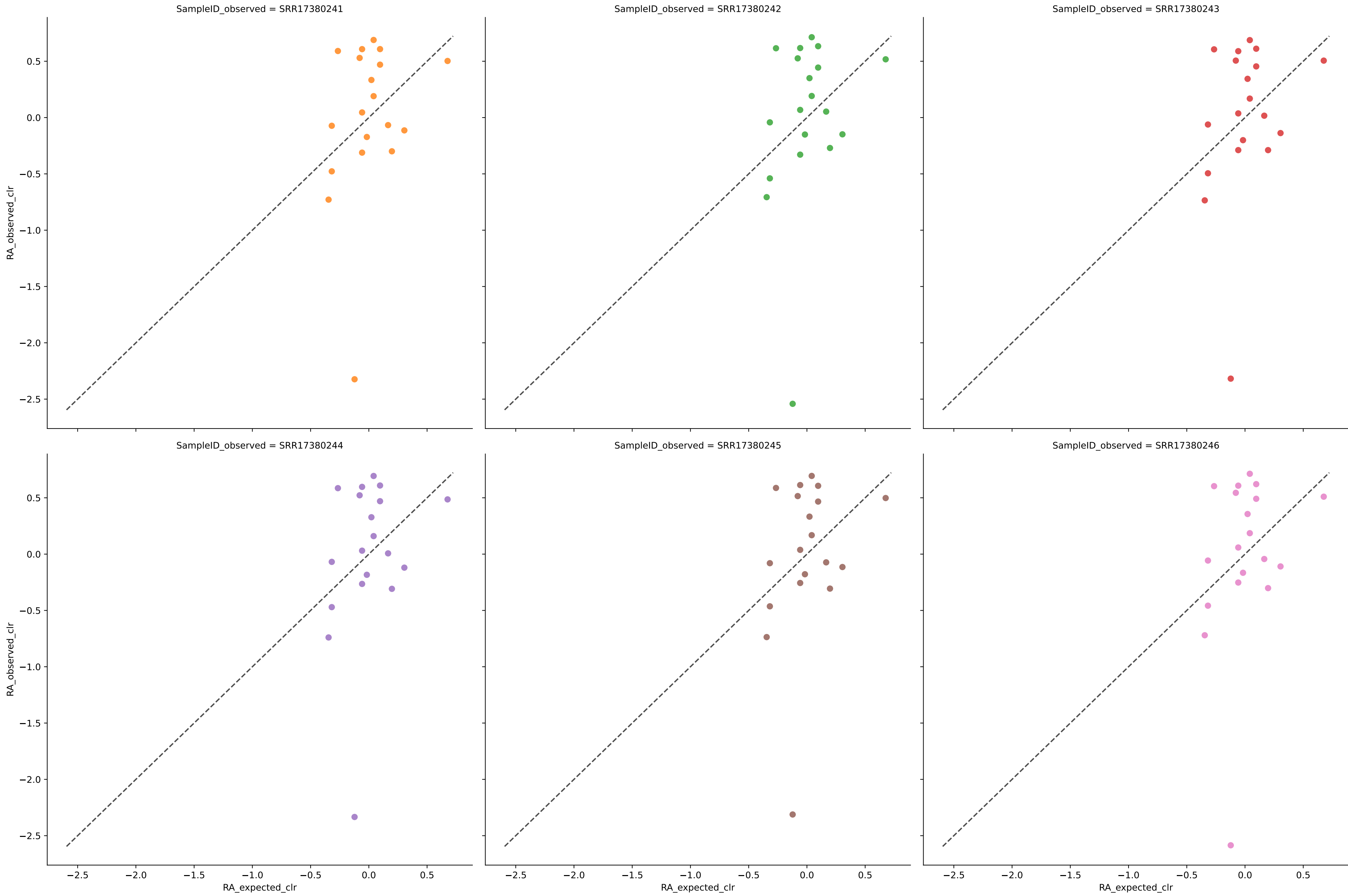
	Diversity	R ²	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR17380241	18	0.0619	0.0067	4.1905	0.6387	0.0082	100.0000	0.0000
SRR17380242	18	0.0655	0.0066	4.1572	0.6414	0.0081	100.0000	0.0000
SRR17380243	18	0.0618	0.0067	4.1904	0.6385	0.0082	100.0000	0.0000
SRR17380244	18	0.0608	0.0067	4.1939	0.6374	0.0083	100.0000	0.0000
SRR17380245	18	0.0609	0.0067	4.2048	0.6373	0.0083	100.0000	0.0000
SRR17380246	18	0.0611	0.0067	4.2050	0.6370	0.0083	100.0000	0.0000
Average	18	0.0620	0.0067	4.1903	0.6384	0.0082	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using bio4 in Experiment tourlousse with filter 0.001



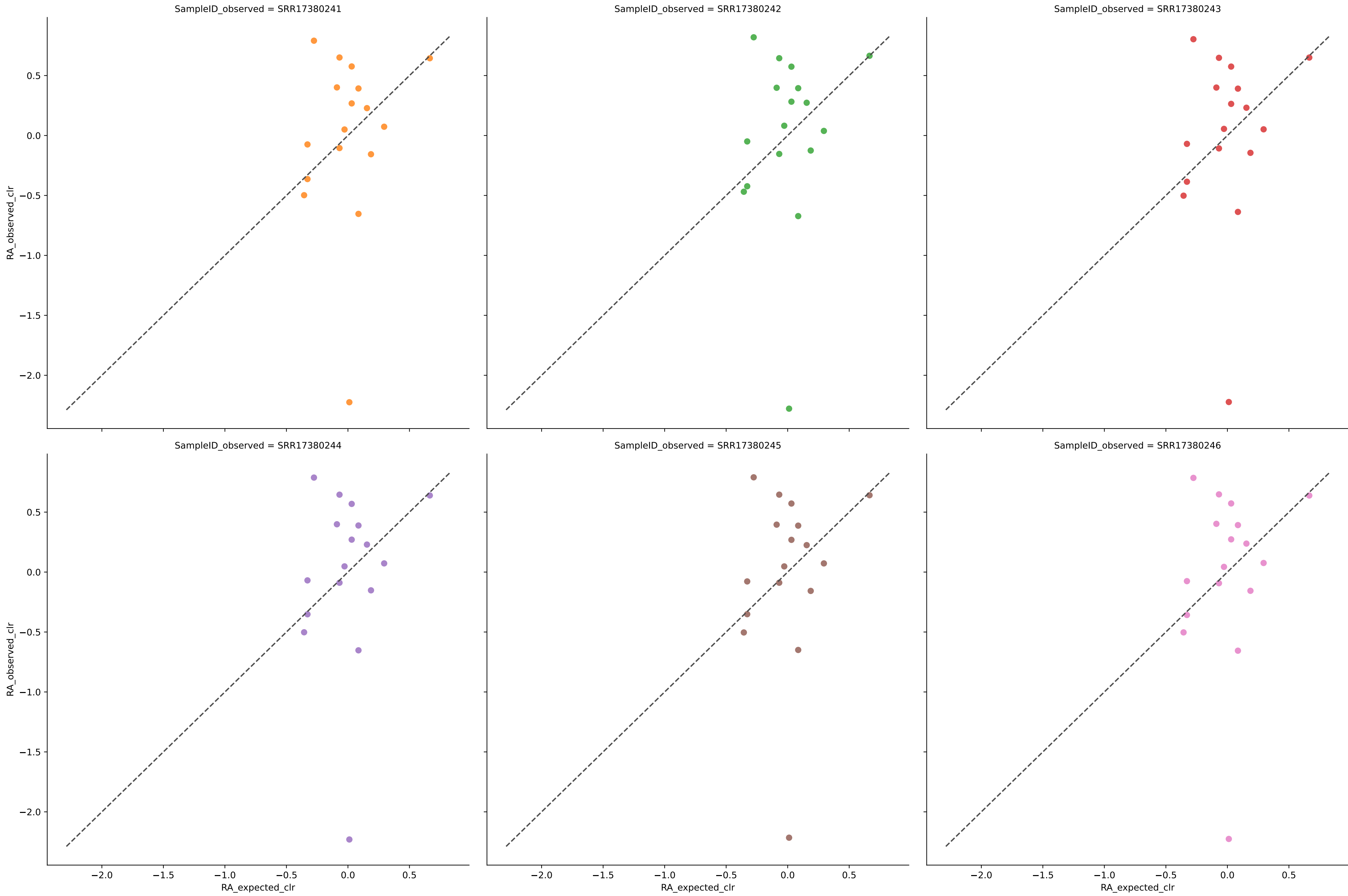
	Diversity	R ²	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR17380241	15	0.6588	0.0018	1.7116	0.9196	0.0023	100.0000	0.0000
SRR17380242	15	0.6349	0.0018	1.8567	0.9185	0.0024	100.0000	0.0000
SRR17380243	15	0.6416	0.0018	1.9887	0.9203	0.0024	100.0000	0.0000
SRR17380244	15	0.6284	0.0019	1.8273	0.9153	0.0024	100.0000	0.0000
SRR17380245	15	0.6673	0.0017	1.6908	0.9221	0.0023	100.0000	0.0000
SRR17380246	15	0.6450	0.0018	1.8257	0.9199	0.0024	100.0000	0.0000
Average	15	0.6460	0.0018	1.8168	0.9193	0.0024	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using jams in Experiment tourlousse with filter 0.001



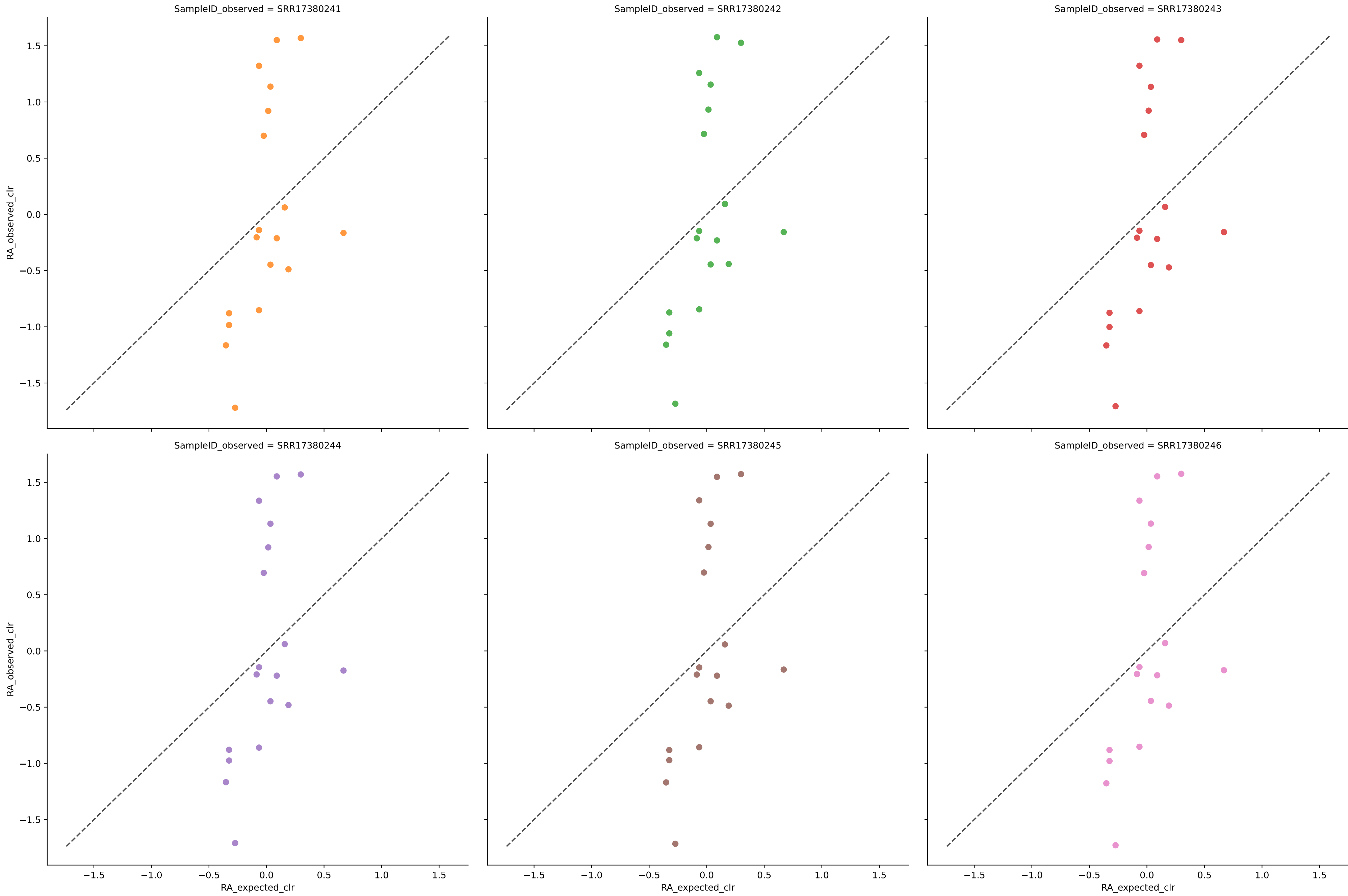
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR17380241	19	0.0685	0.0035	2.8585	0.8028	0.0040	100.0000	0.0000
SRR17380242	19	0.0699	0.0034	3.0443	0.8034	0.0041	100.0000	0.0000
SRR17380243	19	0.0715	0.0034	2.8464	0.8049	0.0040	100.0000	0.0000
SRR17380244	19	0.0671	0.0034	2.8565	0.8044	0.0040	100.0000	0.0000
SRR17380245	19	0.0674	0.0034	2.8435	0.8038	0.0040	100.0000	0.0000
SRR17380246	19	0.0661	0.0035	3.0762	0.8022	0.0041	100.0000	0.0000
Average	19	0.0684	0.0034	2.9209	0.8036	0.0040	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using wgsa in Experiment tourlousse with filter 0.001



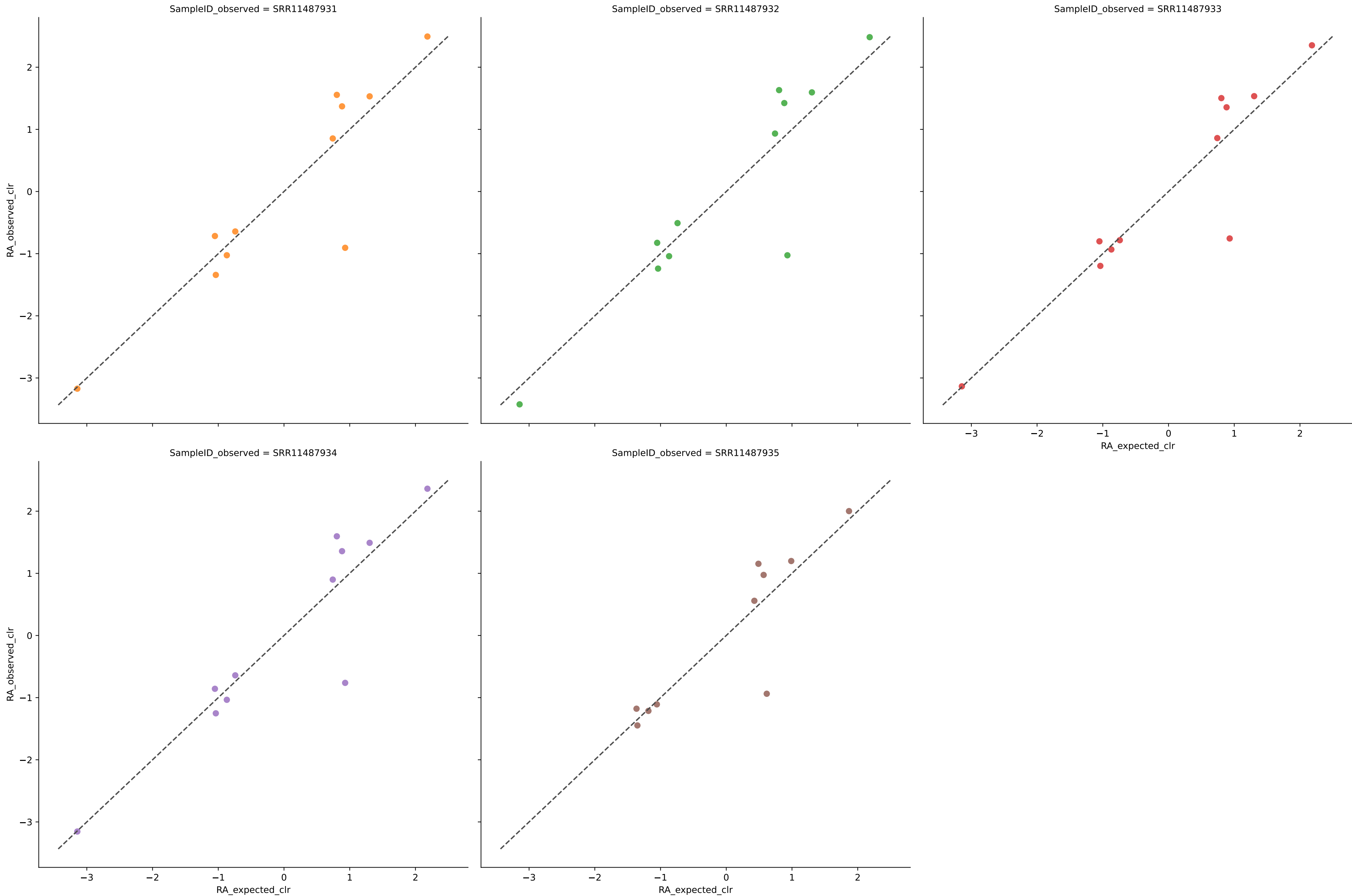
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR17380241	17	0.0588	0.0036	2.8562	0.8157	0.0047	100.0000	0.0000
SRR17380242	17	0.0603	0.0036	2.9180	0.8144	0.0047	100.0000	0.0000
SRR17380243	17	0.0586	0.0036	2.8552	0.8154	0.0047	100.0000	0.0000
SRR17380244	17	0.0579	0.0036	2.8562	0.8166	0.0047	100.0000	0.0000
SRR17380245	17	0.0583	0.0036	2.8440	0.8167	0.0047	100.0000	0.0000
SRR17380246	17	0.0586	0.0036	2.8549	0.8162	0.0047	100.0000	0.0000
Average	17	0.0587	0.0036	2.8641	0.8158	0.0047	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using woltka in Experiment tourlousse with filter 0.001



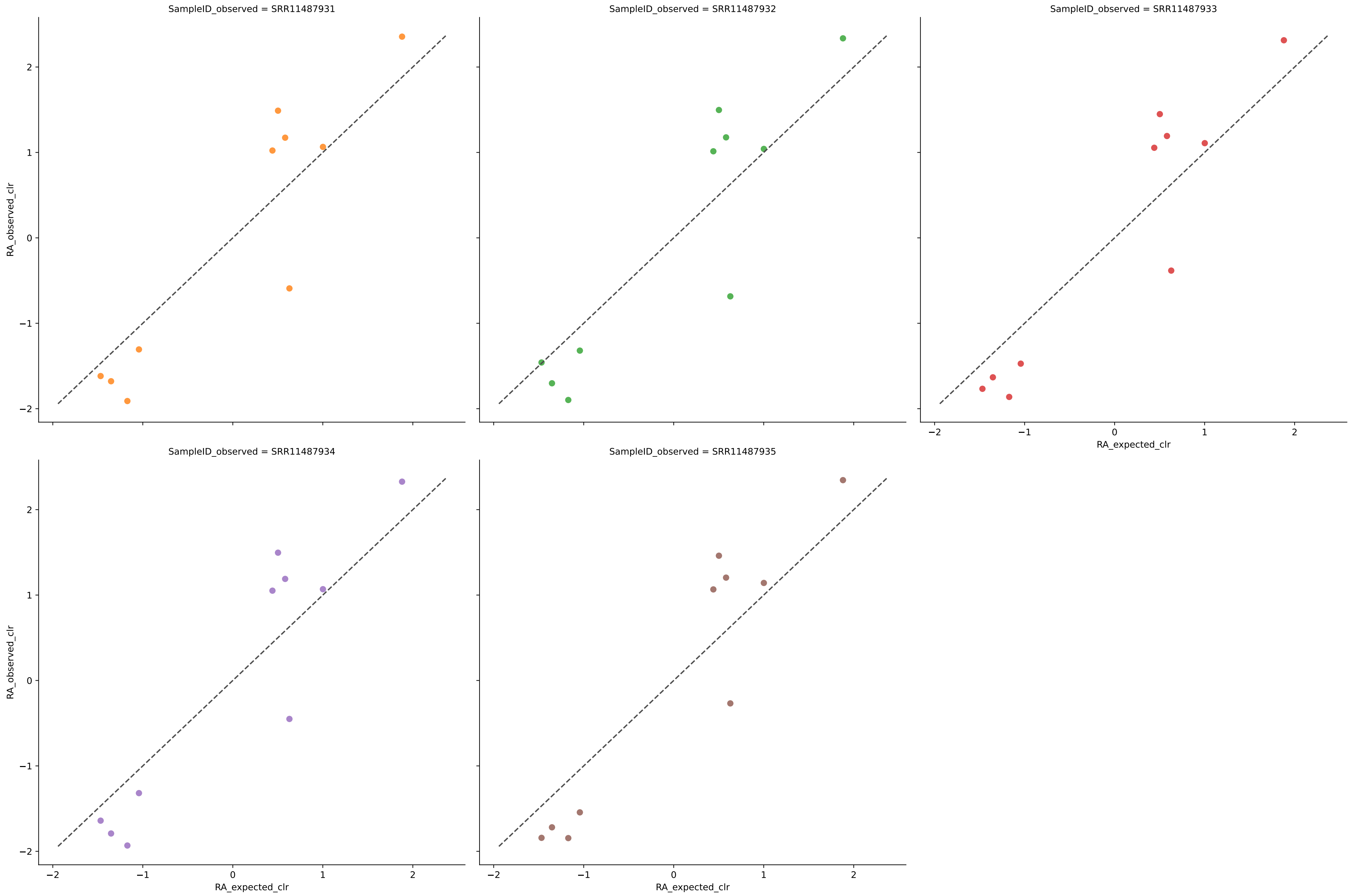
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR17380241	18	0.0560	0.0073	3.7226	0.6054	0.0086	100.0000	0.0000
SRR17380242	18	0.0568	0.0072	3.6942	0.6082	0.0084	100.0000	0.0000
SRR17380243	18	0.0555	0.0073	3.7162	0.6058	0.0086	100.0000	0.0000
SRR17380244	18	0.0550	0.0073	3.7255	0.6044	0.0086	100.0000	0.0000
SRR17380245	18	0.0554	0.0073	3.7281	0.6042	0.0086	100.0000	0.0000
SRR17380246	18	0.0555	0.0073	3.7368	0.6044	0.0086	100.0000	0.0000
Average	18	0.0557	0.0073	3.7206	0.6054	0.0086	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using bio4 in Experiment Amos hilo with filter 0.001



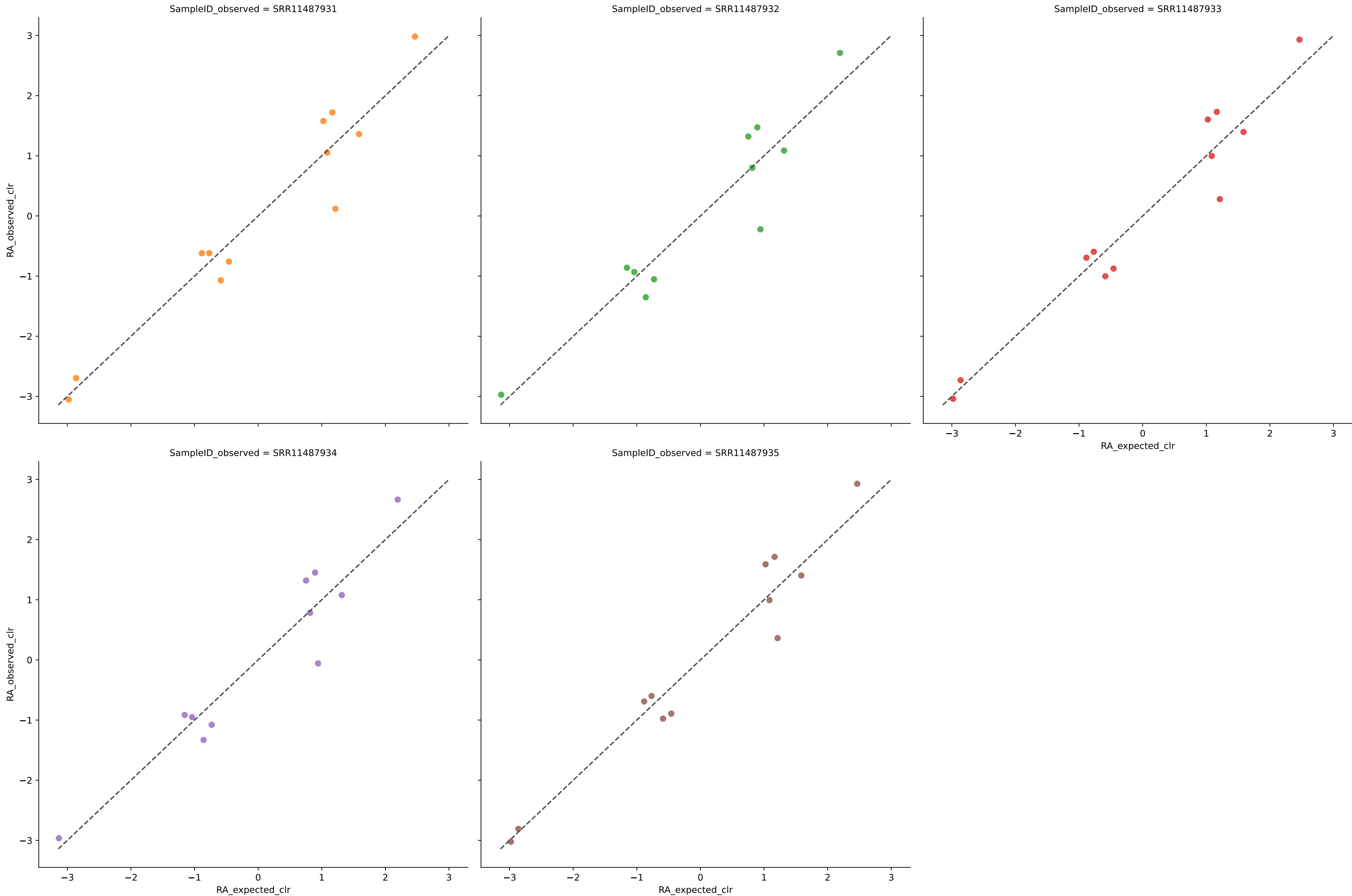
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487931	11	0.9035	0.0044	2.1389	0.8778	0.0073	100.0000	0.0000
SRR11487932	11	0.8892	0.0043	2.2959	0.8825	0.0075	100.0000	0.0000
SRR11487933	11	0.8928	0.0040	1.9349	0.8892	0.0071	100.0000	0.0000
SRR11487934	11	0.8791	0.0041	1.9826	0.8879	0.0076	100.0000	0.0000
SRR11487935	10	0.8906	0.0042	1.7737	0.8948	0.0072	100.0000	0.0000
Average	11	0.8910	0.0042	2.0252	0.8864	0.0074	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using jams in Experiment Amos hilo with filter 0.001



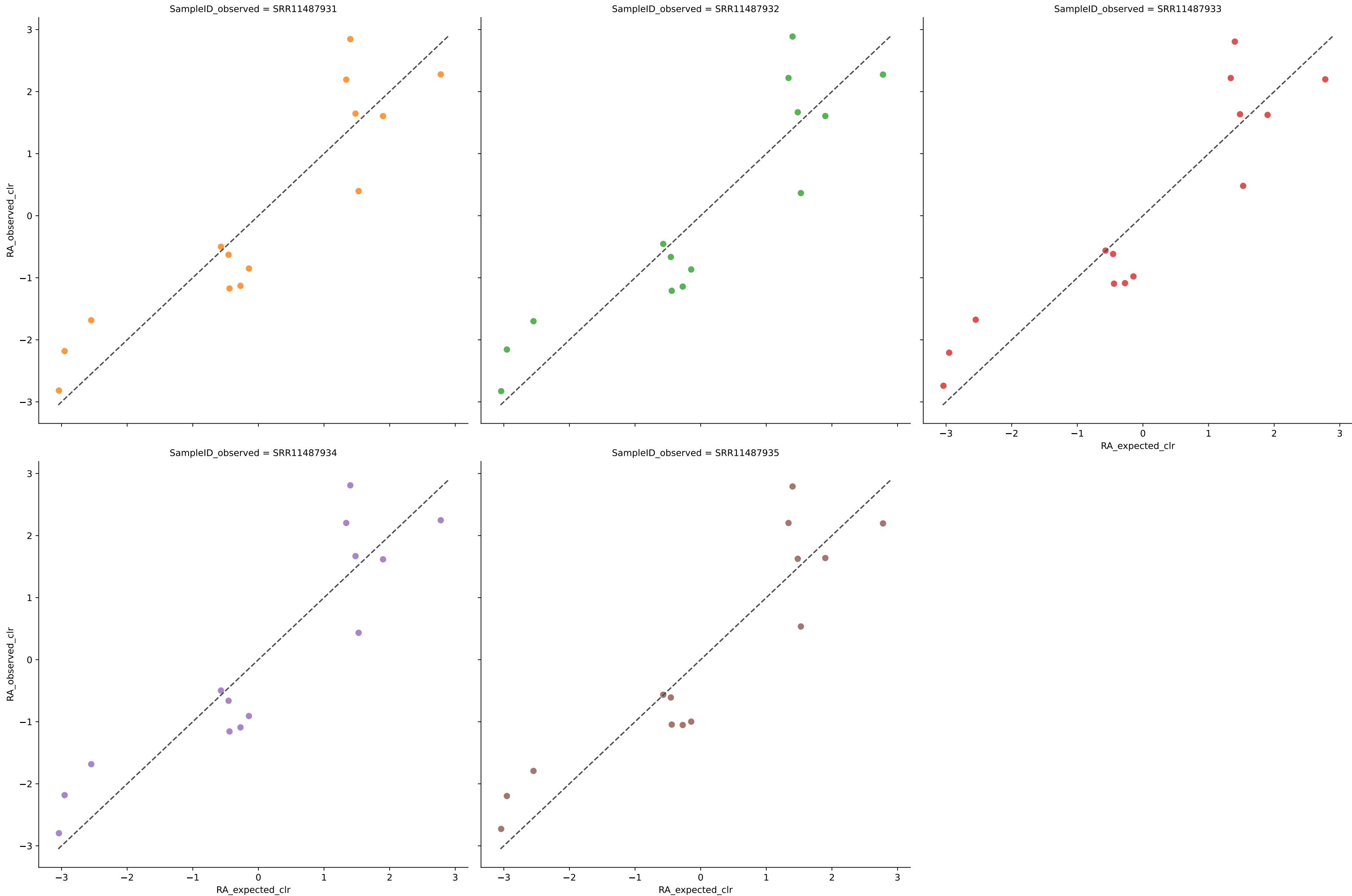
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487931	10	0.8808	0.0065	2.0309	0.8364	0.0086	100.0000	0.0000
SRR11487932	10	0.8724	0.0066	2.0817	0.8345	0.0088	100.0000	0.0000
SRR11487933	10	0.8884	0.0062	1.9233	0.8462	0.0080	100.0000	0.0000
SRR11487934	10	0.8777	0.0064	1.9941	0.8392	0.0084	100.0000	0.0000
SRR11487935	10	0.8961	0.0061	1.9280	0.8479	0.0078	100.0000	0.0000
Average	10	0.8831	0.0064	1.9916	0.8408	0.0083	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using wgsa in Experiment Amos hilo with filter 0.001



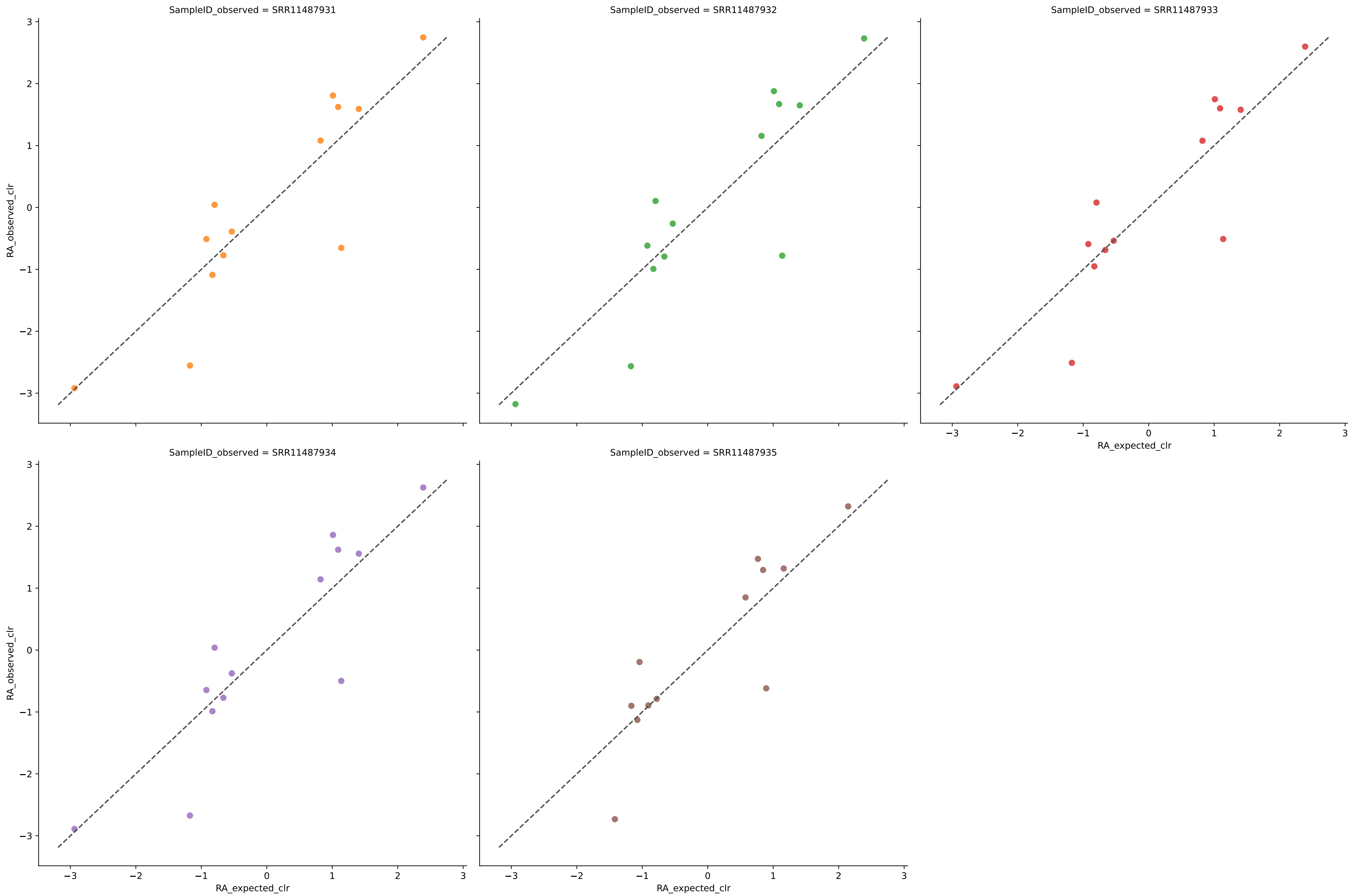
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487931	12	0.9082	0.0061	1.6060	0.8169	0.0094	100.0000	0.0000
SRR11487932	11	0.9042	0.0067	1.6737	0.8153	0.0097	100.0000	0.0000
SRR11487933	12	0.9141	0.0059	1.4896	0.8244	0.0086	100.0000	0.0000
SRR11487934	11	0.9089	0.0065	1.5331	0.8223	0.0092	100.0000	0.0000
SRR11487935	12	0.9181	0.0057	1.4180	0.8282	0.0084	100.0000	0.0000
Average	12	0.9107	0.0062	1.5441	0.8214	0.0091	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using wol in Experiment Amos hilo with filter 0.001



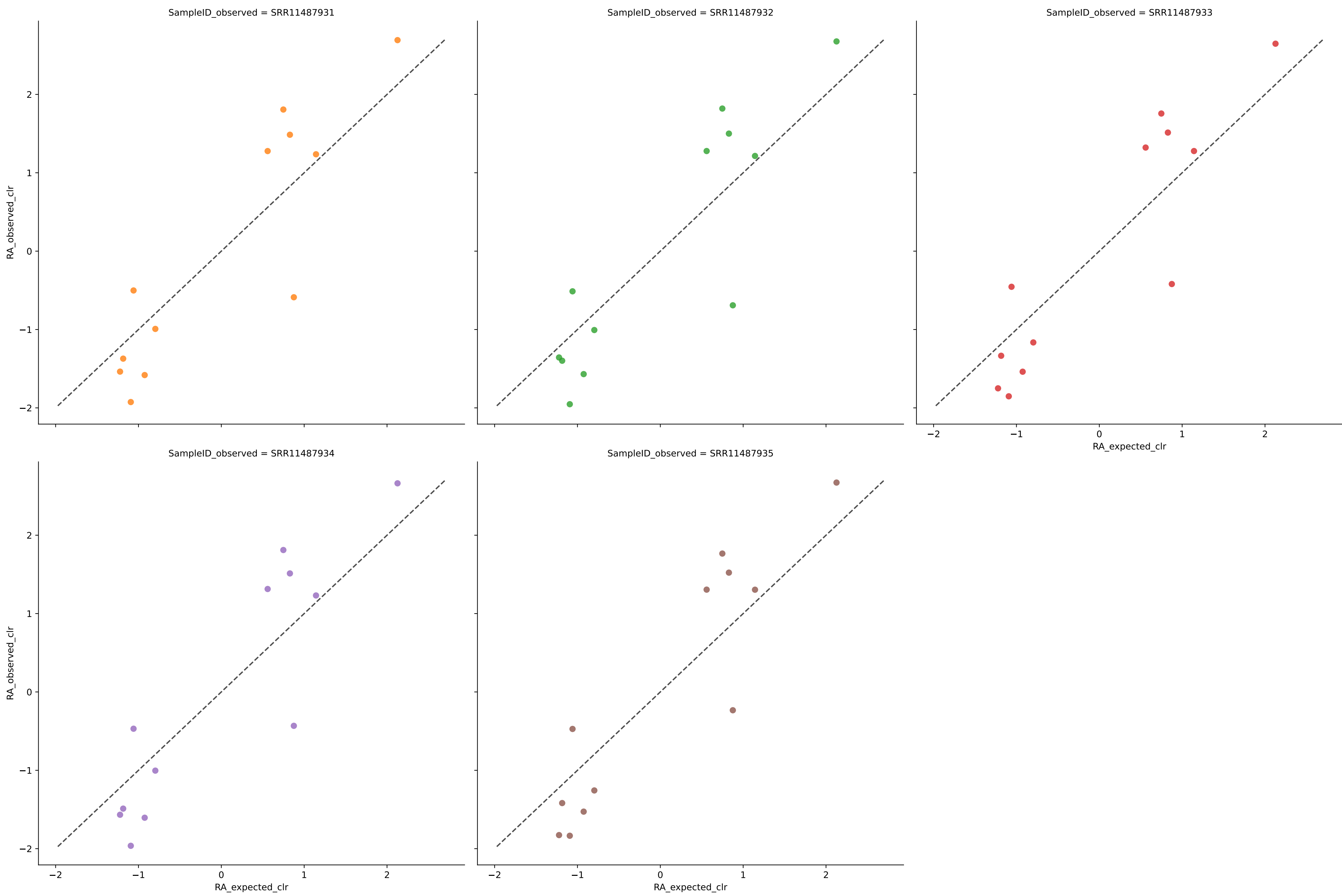
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487931	14	0.3425	0.0099	2.7670	0.6546	0.0178	100.0000	0.0000
SRR11487932	14	0.3234	0.0101	2.8351	0.6457	0.0183	100.0000	0.0000
SRR11487933	14	0.3281	0.0100	2.7385	0.6511	0.0180	100.0000	0.0000
SRR11487934	14	0.3458	0.0099	2.7456	0.6552	0.0177	100.0000	0.0000
SRR11487935	14	0.3355	0.0098	2.6575	0.6566	0.0178	100.0000	0.0000
Average	14	0.3351	0.0099	2.7487	0.6526	0.0179	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using bio4 in Experiment Amos hilo with filter 0.001



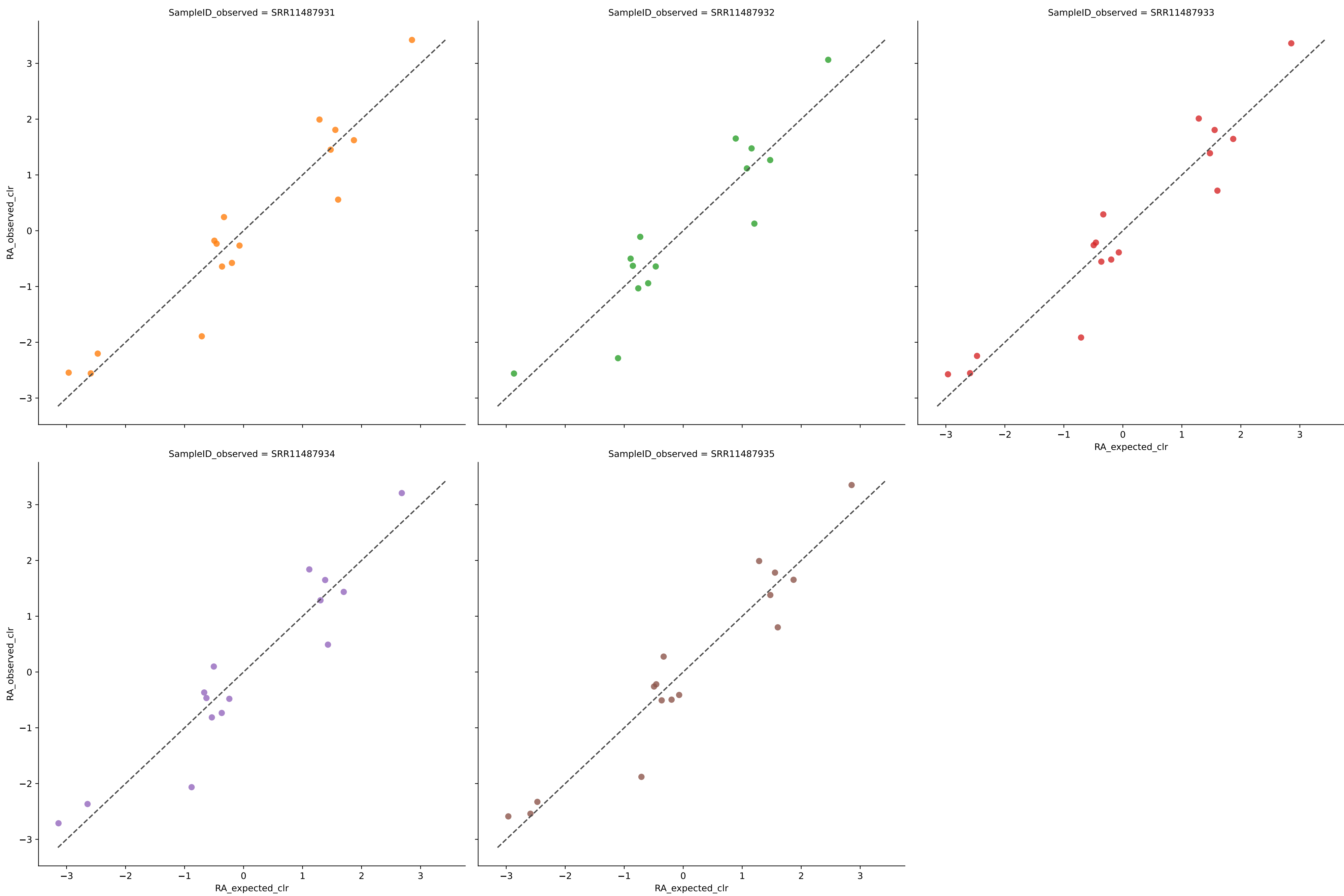
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487931	13	0.9051	0.0041	2.6892	0.8660	0.0068	100.0000	0.0000
SRR11487932	13	0.8918	0.0039	2.8399	0.8717	0.0070	100.0000	0.0000
SRR11487933	13	0.8956	0.0037	2.5177	0.8796	0.0066	100.0000	0.0000
SRR11487934	13	0.8820	0.0040	2.6358	0.8693	0.0071	100.0000	0.0000
SRR11487935	12	0.8960	0.0039	2.3734	0.8830	0.0067	100.0000	0.0000
Average	13	0.8941	0.0039	2.6112	0.8739	0.0068	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using jams in Experiment Amos hilo with filter 0.001



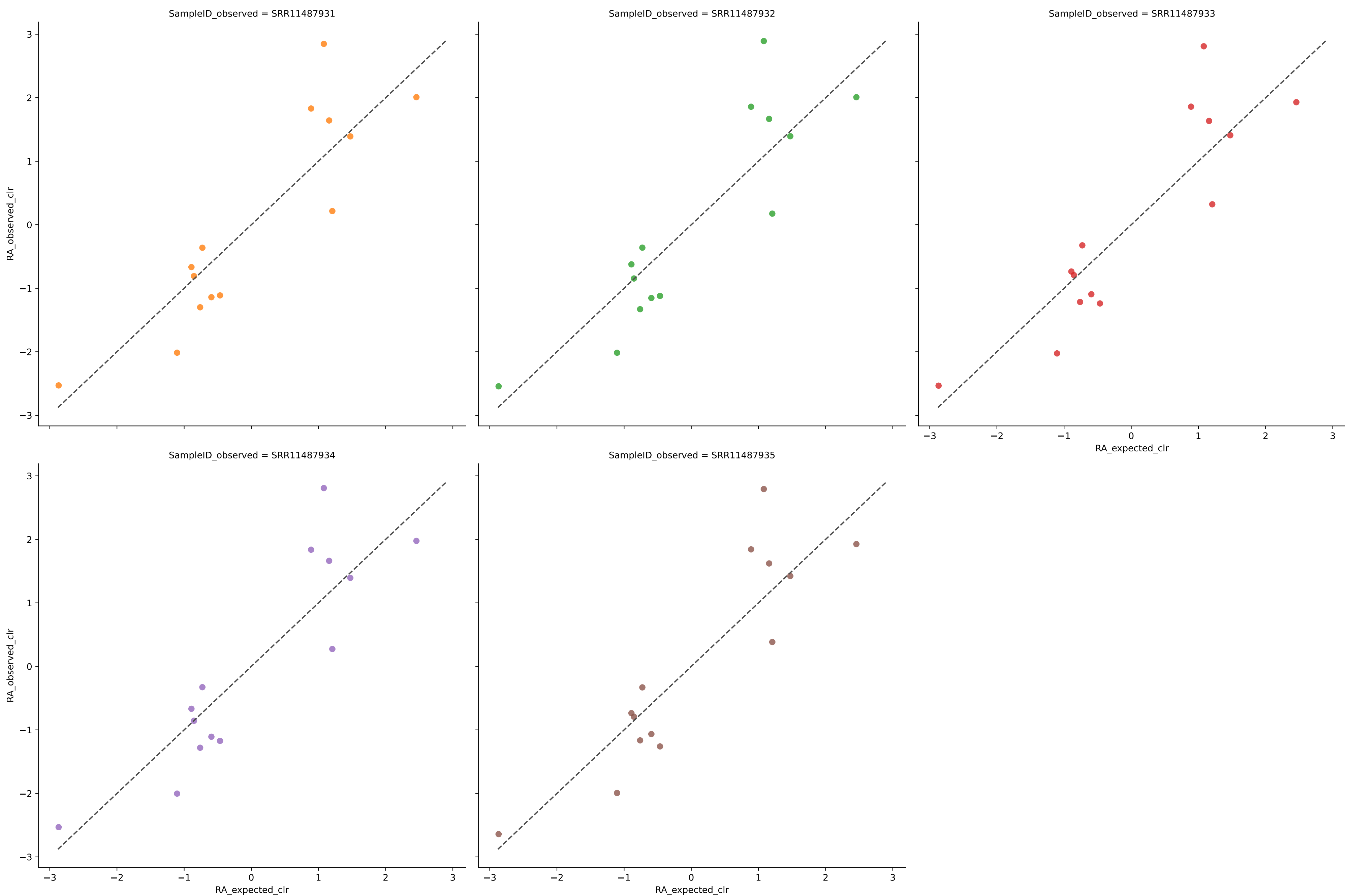
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487931	12	0.8855	0.0060	2.4794	0.8201	0.0083	100.0000	0.0000
SRR11487932	12	0.8779	0.0061	2.5372	0.8176	0.0085	100.0000	0.0000
SRR11487933	12	0.8897	0.0058	2.4059	0.8273	0.0078	100.0000	0.0000
SRR11487934	12	0.8812	0.0059	2.4483	0.8215	0.0081	100.0000	0.0000
SRR11487935	12	0.8978	0.0057	2.3490	0.8302	0.0076	100.0000	0.0000
Average	12	0.8864	0.0059	2.4440	0.8234	0.0081	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using wgsa in Experiment Amos hilo with filter 0.001



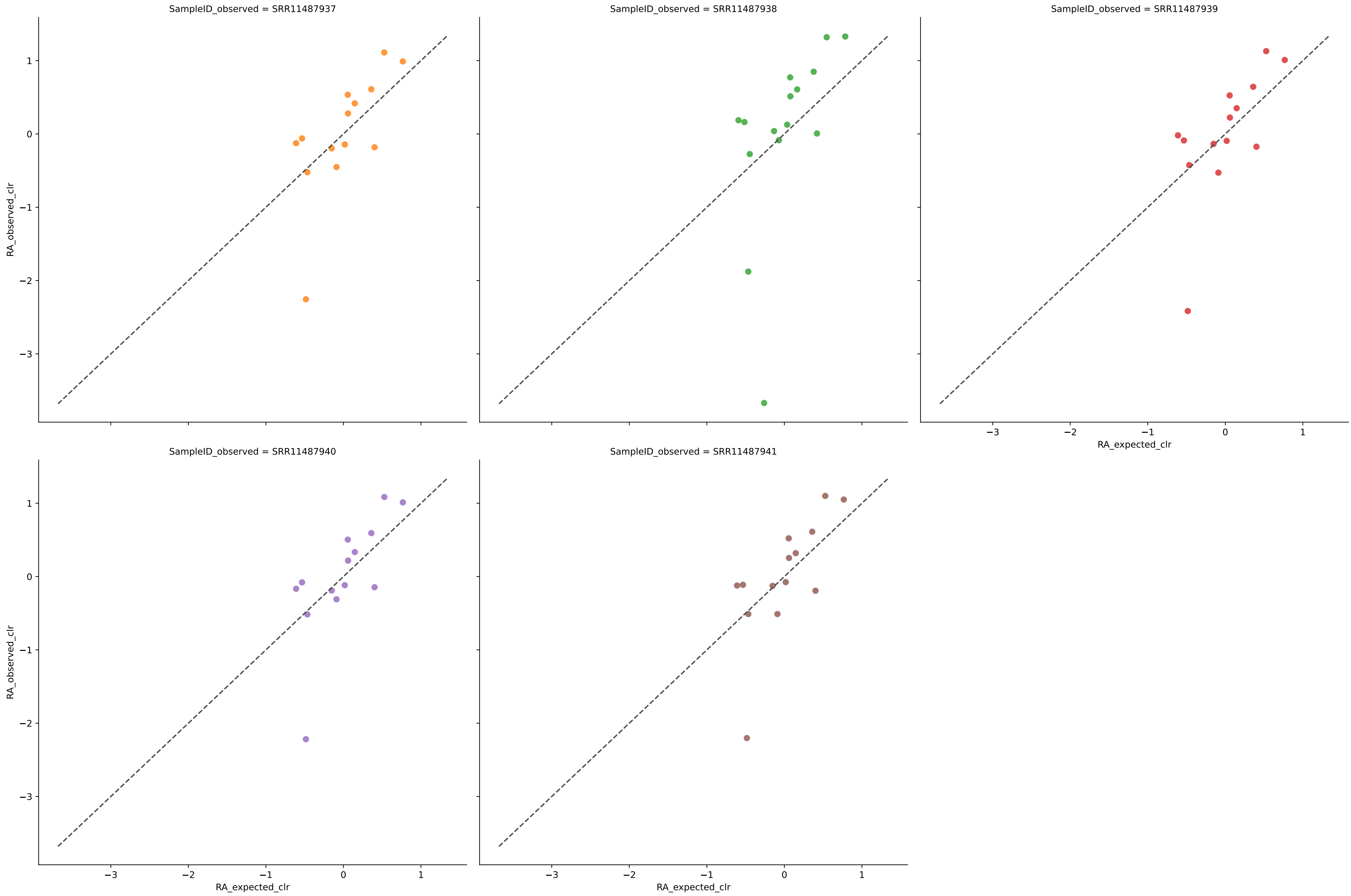
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487931	16	0.9115	0.0047	2.1050	0.8138	0.0087	100.0000	0.0000
SRR11487932	14	0.9094	0.0053	2.1347	0.8123	0.0092	100.0000	0.0000
SRR11487933	16	0.9178	0.0044	2.0224	0.8225	0.0079	100.0000	0.0000
SRR11487934	15	0.9144	0.0048	2.0568	0.8187	0.0084	100.0000	0.0000
SRR11487935	16	0.9215	0.0043	1.9332	0.8271	0.0078	100.0000	0.0000
Average	15	0.9149	0.0047	2.0504	0.8189	0.0084	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using wol in Experiment Amos hilo with filter 0.001



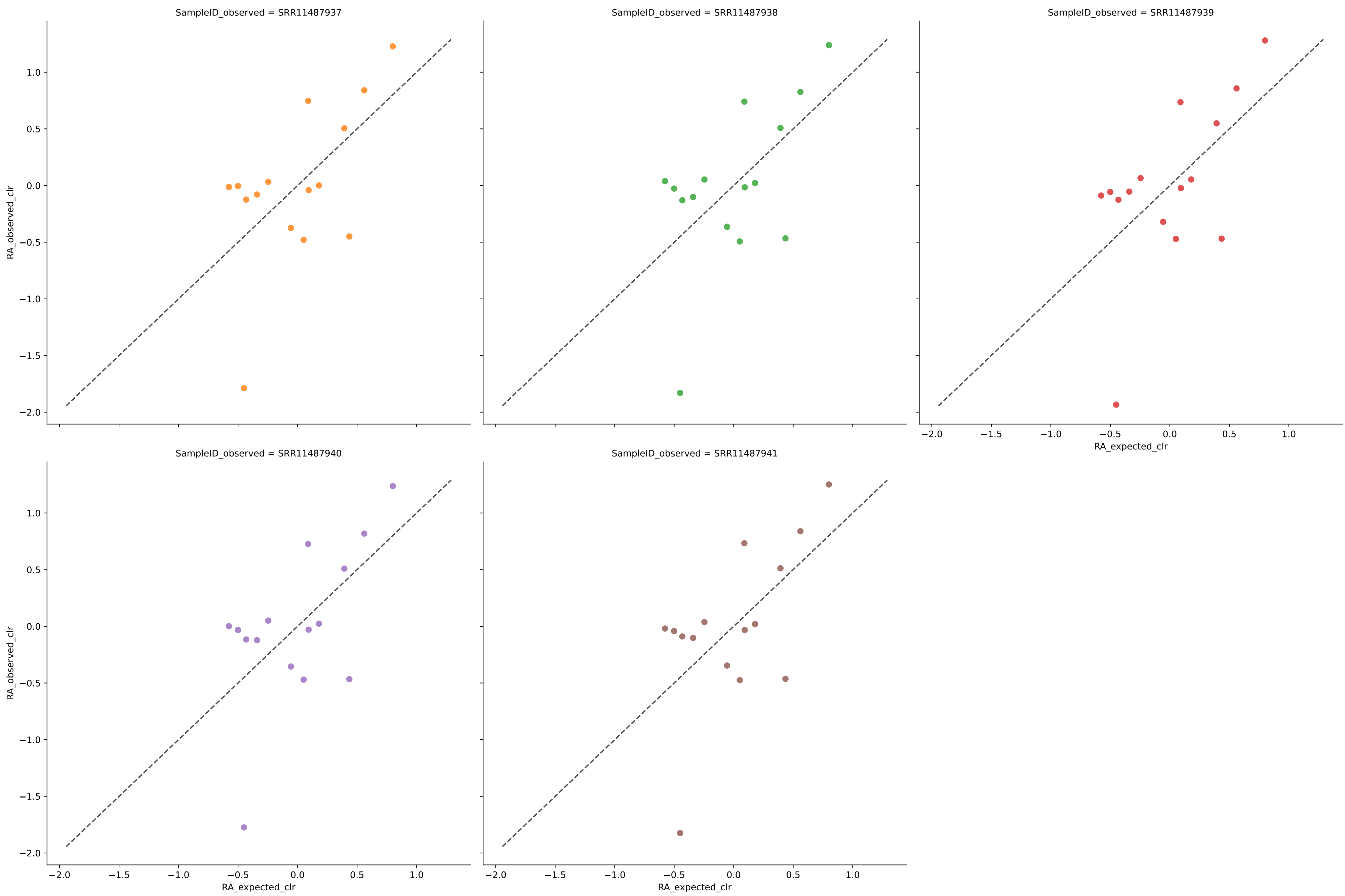
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487931	14	0.2393	0.0107	2.7515	0.6243	0.0202	100.0000	0.0000
SRR11487932	14	0.2243	0.0110	2.8209	0.6154	0.0207	100.0000	0.0000
SRR11487933	14	0.2286	0.0108	2.7261	0.6220	0.0202	100.0000	0.0000
SRR11487934	14	0.2432	0.0107	2.7218	0.6256	0.0199	100.0000	0.0000
SRR11487935	14	0.2348	0.0106	2.6569	0.6279	0.0200	100.0000	0.0000
Average	14	0.2340	0.0108	2.7354	0.6230	0.0202	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using bio4 in Experiment Amos mixed with filter 0.001



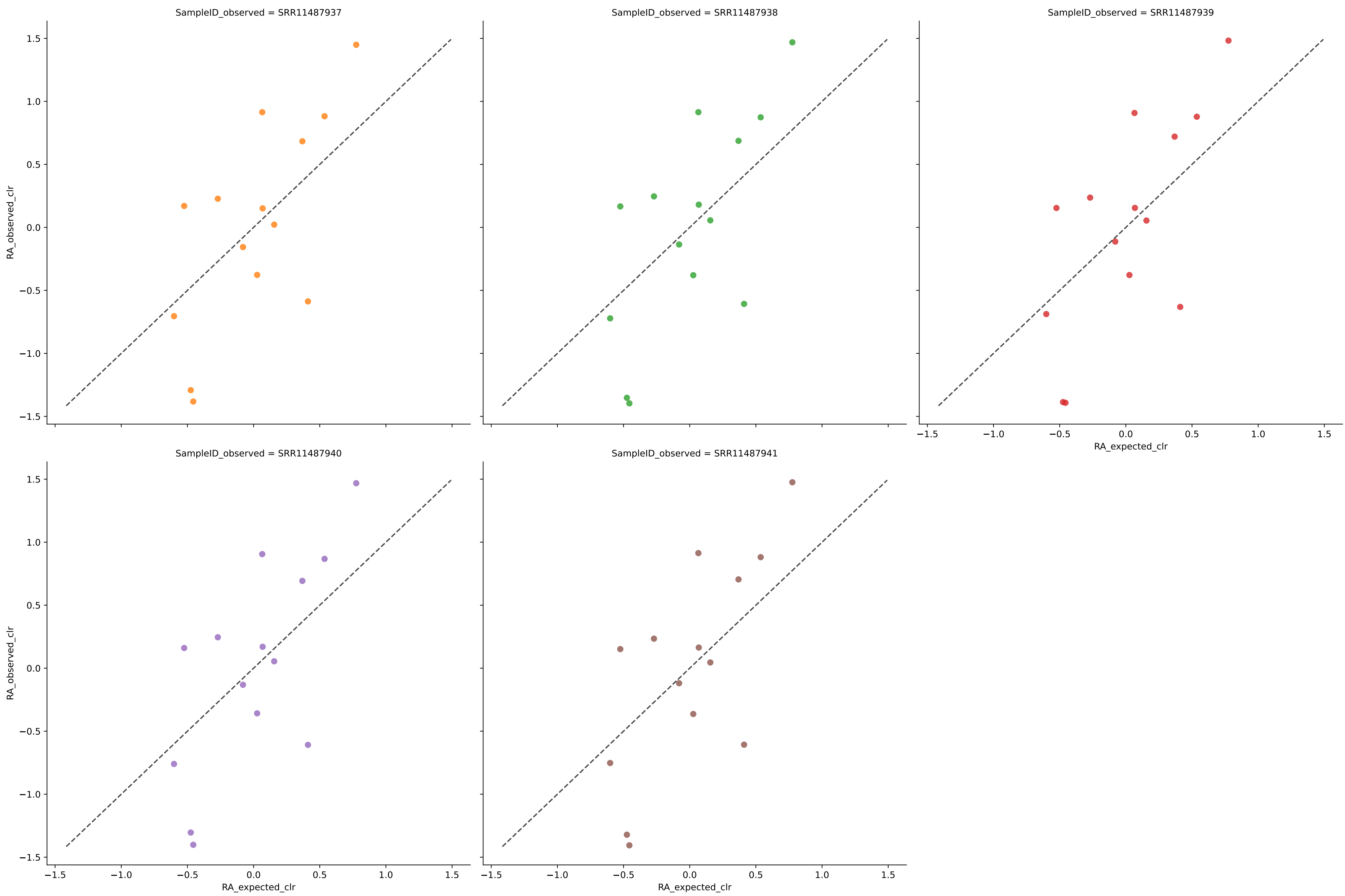
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487937	14	0.6657	0.0042	2.2146	0.8512	0.0053	100.0000	0.0000
SRR11487938	15	0.6856	0.0045	4.1118	0.8333	0.0056	100.0000	0.0000
SRR11487939	14	0.6600	0.0041	2.3682	0.8548	0.0054	100.0000	0.0000
SRR11487940	14	0.7007	0.0039	2.1144	0.8628	0.0050	100.0000	0.0000
SRR11487941	14	0.6936	0.0041	2.1610	0.8562	0.0053	100.0000	0.0000
Average	14	0.6811	0.0042	2.5940	0.8517	0.0053	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using jams in Experiment Amos mixed with filter 0.001



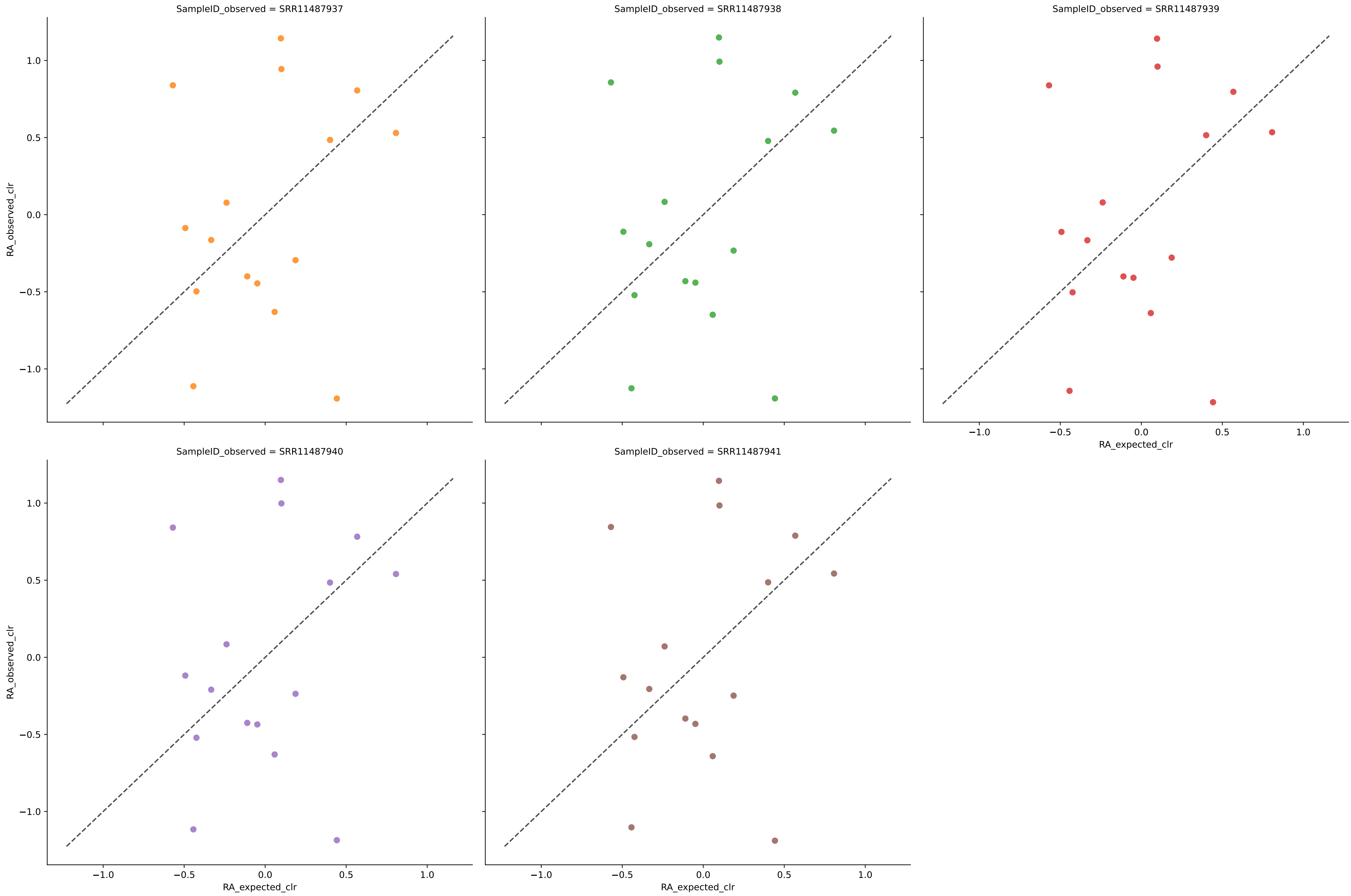
	Diversity	R ²	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487937	15	0.5720	0.0047	2.1252	0.8246	0.0057	100.0000	0.0000
SRR11487938	15	0.5697	0.0047	2.1639	0.8250	0.0058	100.0000	0.0000
SRR11487939	15	0.5984	0.0046	2.2036	0.8278	0.0057	100.0000	0.0000
SRR11487940	15	0.5777	0.0046	2.1051	0.8269	0.0057	100.0000	0.0000
SRR11487941	15	0.5820	0.0046	2.1406	0.8259	0.0058	100.0000	0.0000
Average	15	0.5800	0.0046	2.1477	0.8260	0.0058	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using wgsa in Experiment Amos mixed with filter 0.001



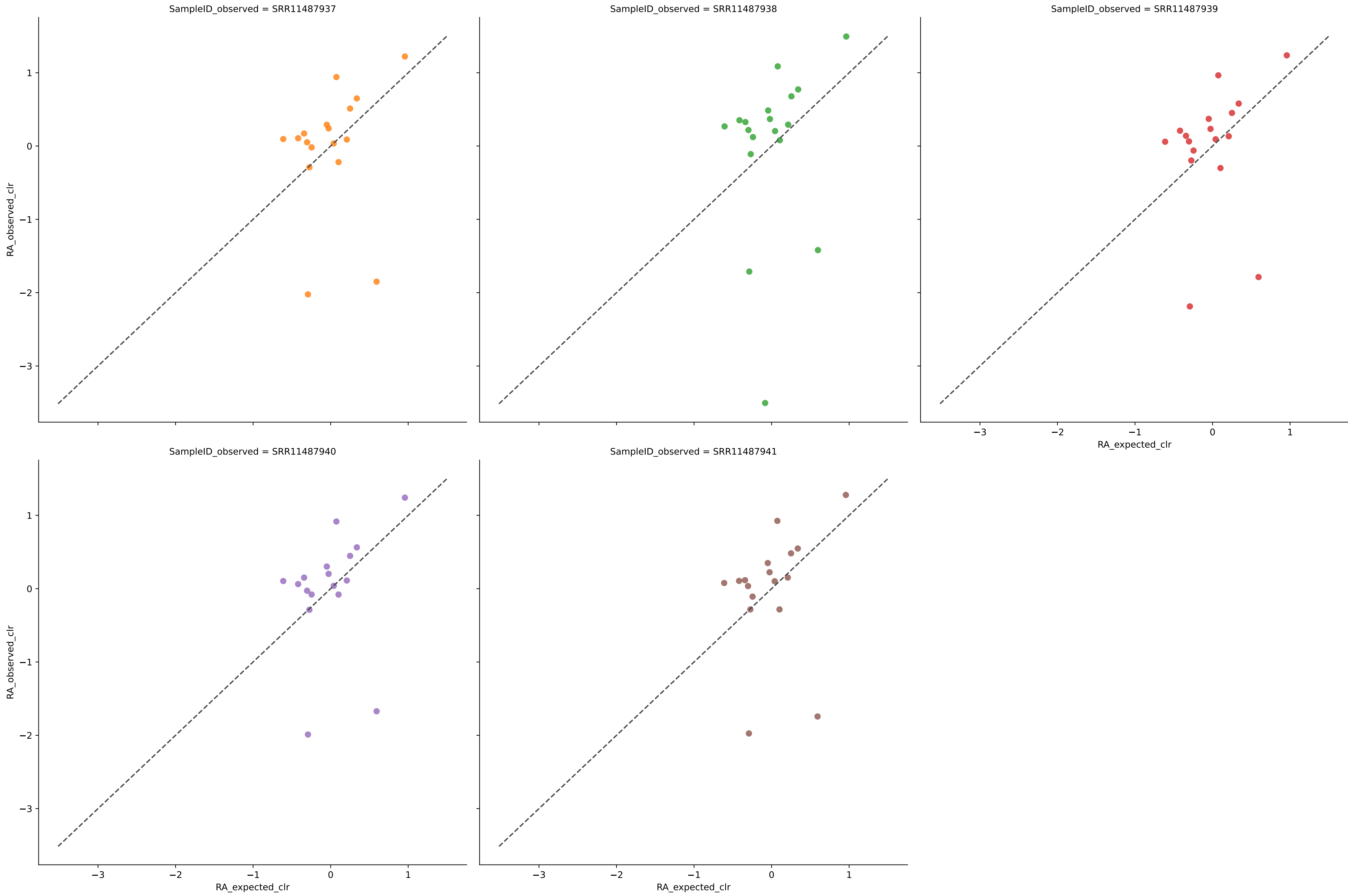
	Diversity	R ²	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487937	14	0.6079	0.0060	2.2021	0.7890	0.0076	100.0000	0.0000
SRR11487938	14	0.6085	0.0060	2.2487	0.7891	0.0077	100.0000	0.0000
SRR11487939	14	0.6129	0.0061	2.2696	0.7872	0.0077	100.0000	0.0000
SRR11487940	14	0.6117	0.0060	2.2255	0.7896	0.0076	100.0000	0.0000
SRR11487941	14	0.6147	0.0061	2.2373	0.7880	0.0077	100.0000	0.0000
Average	14	0.6112	0.0060	2.2366	0.7886	0.0077	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using wol in Experiment Amos mixed with filter 0.001



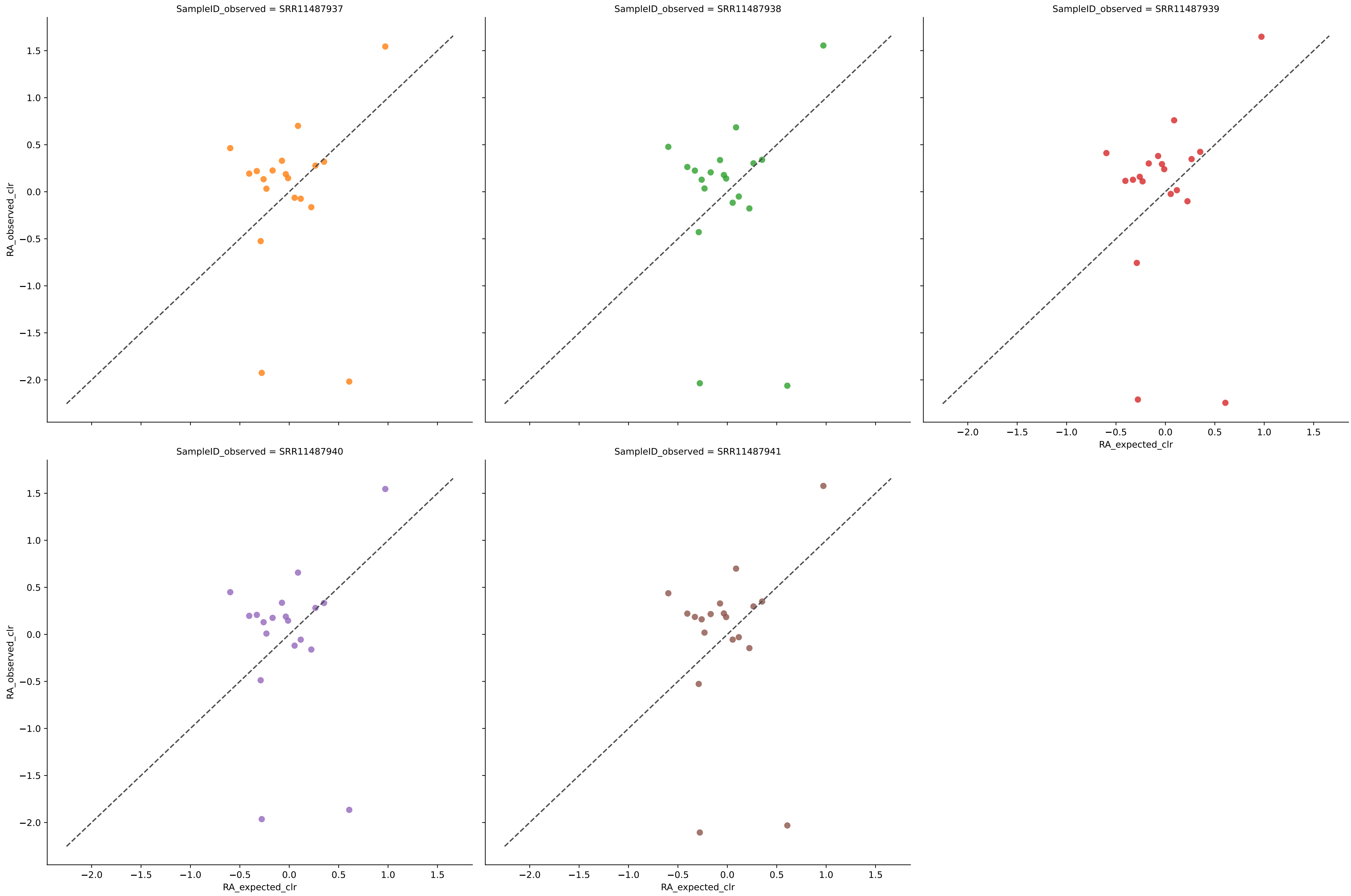
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487937	16	0.0645	0.0066	2.8809	0.7364	0.0087	100.0000	0.0000
SRR11487938	16	0.0633	0.0066	2.9006	0.7340	0.0089	100.0000	0.0000
SRR11487939	16	0.0662	0.0065	2.8965	0.7387	0.0087	100.0000	0.0000
SRR11487940	16	0.0643	0.0066	2.8819	0.7351	0.0088	100.0000	0.0000
SRR11487941	16	0.0658	0.0066	2.8748	0.7369	0.0088	100.0000	0.0000
Average	16	0.0648	0.0066	2.8869	0.7362	0.0088	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using bio4 in Experiment Amos mixed with filter 0.001



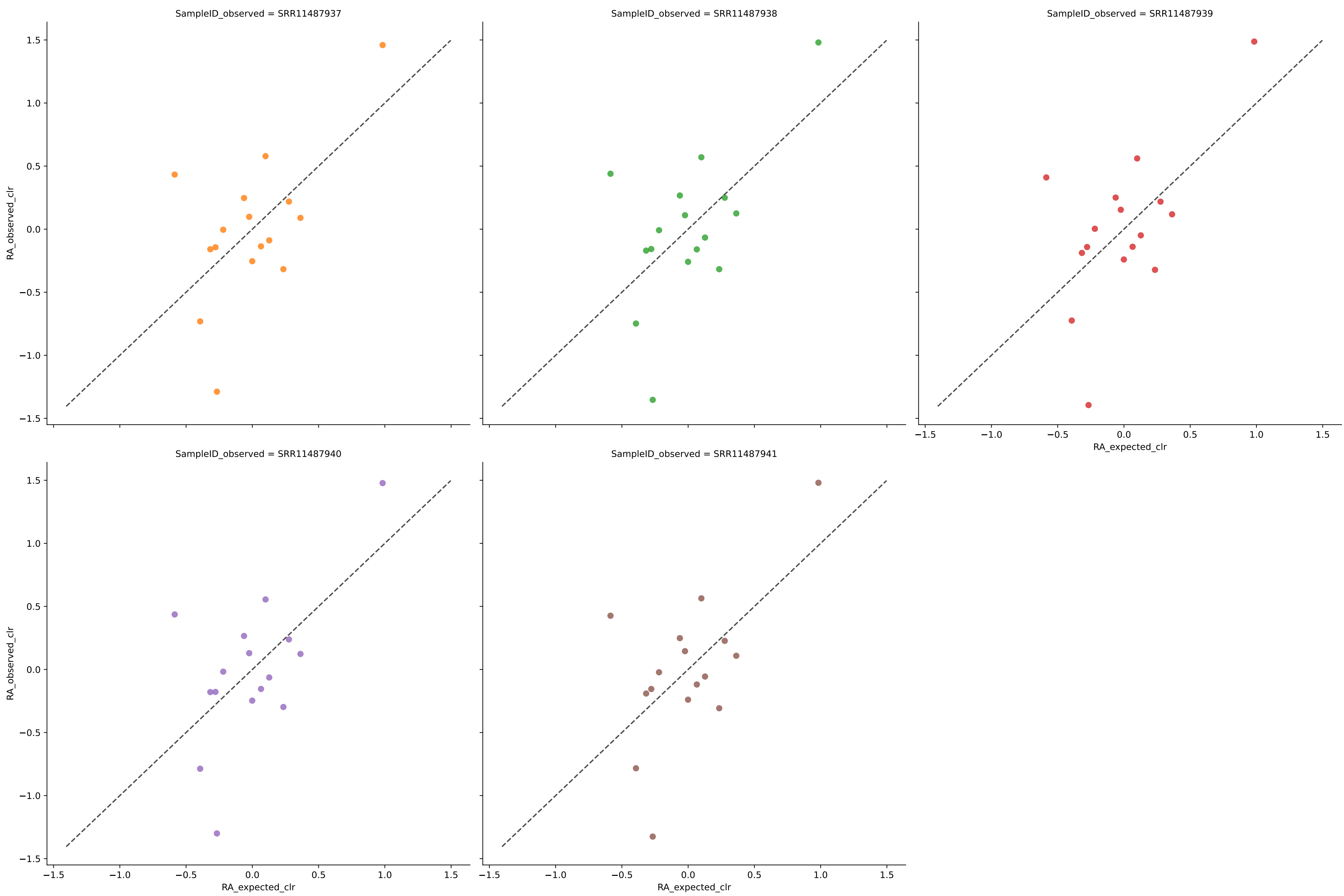
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487937	17	0.3104	0.0042	3.3865	0.8188	0.0061	100.0000	0.0000
SRR11487938	18	0.3547	0.0044	4.7066	0.8048	0.0062	100.0000	0.0000
SRR11487939	17	0.3031	0.0041	3.4478	0.8228	0.0061	100.0000	0.0000
SRR11487940	17	0.3409	0.0040	3.1931	0.8304	0.0059	100.0000	0.0000
SRR11487941	17	0.3459	0.0041	3.2683	0.8246	0.0060	100.0000	0.0000
Average	17	0.3310	0.0042	3.6005	0.8203	0.0061	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using jams in Experiment Amos mixed with filter 0.001



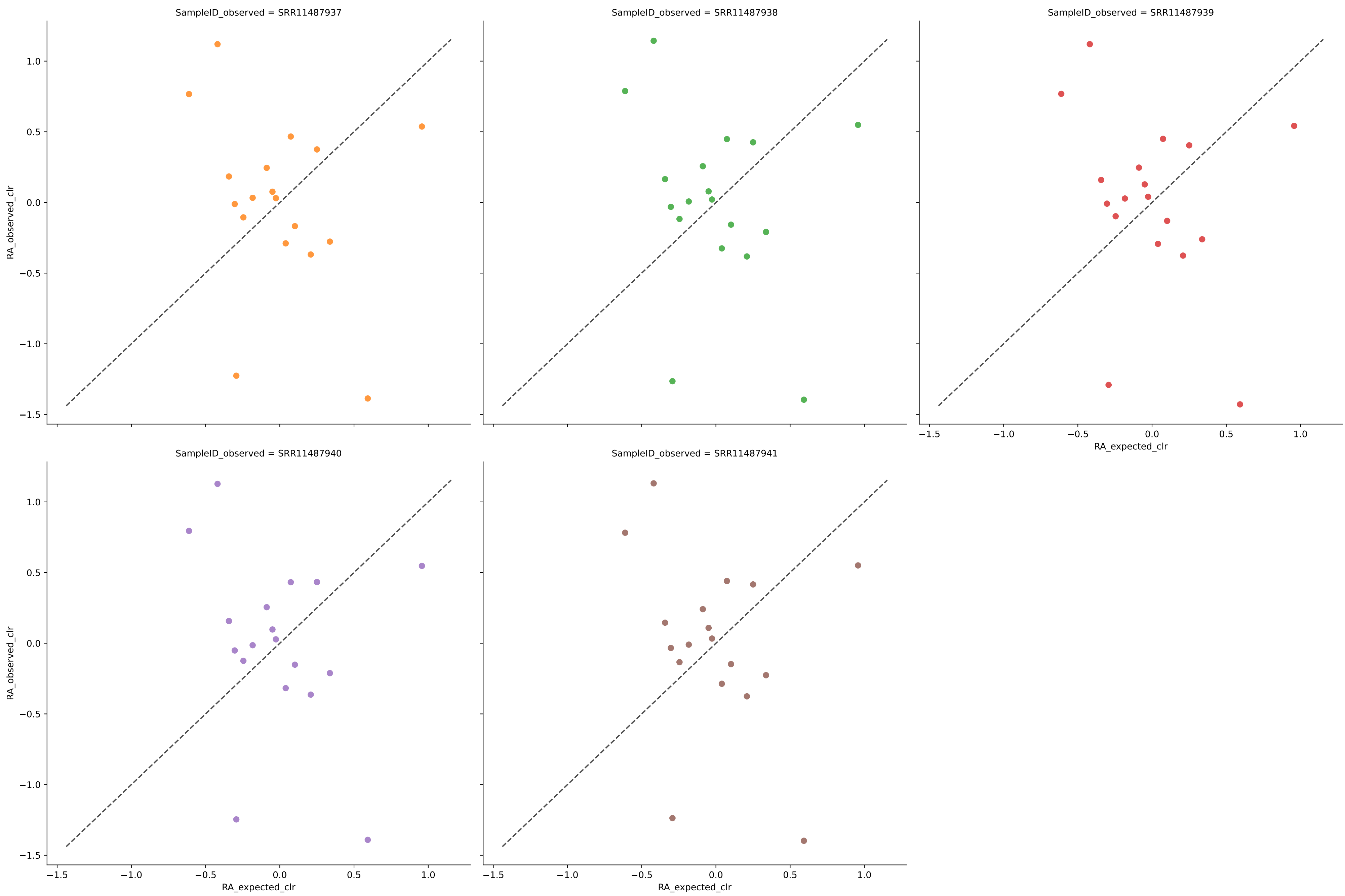
	Diversity	R ²	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487937	19	0.3708	0.0043	3.6003	0.7947	0.0060	100.0000	0.0000
SRR11487938	19	0.3682	0.0044	3.6950	0.7941	0.0061	100.0000	0.0000
SRR11487939	19	0.4162	0.0042	3.9369	0.7974	0.0060	100.0000	0.0000
SRR11487940	19	0.3834	0.0043	3.4885	0.7968	0.0060	100.0000	0.0000
SRR11487941	19	0.3887	0.0043	3.6944	0.7964	0.0060	100.0000	0.0000
Average	19	0.3854	0.0043	3.6830	0.7959	0.0060	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using wgsa in Experiment Amos mixed with filter 0.001



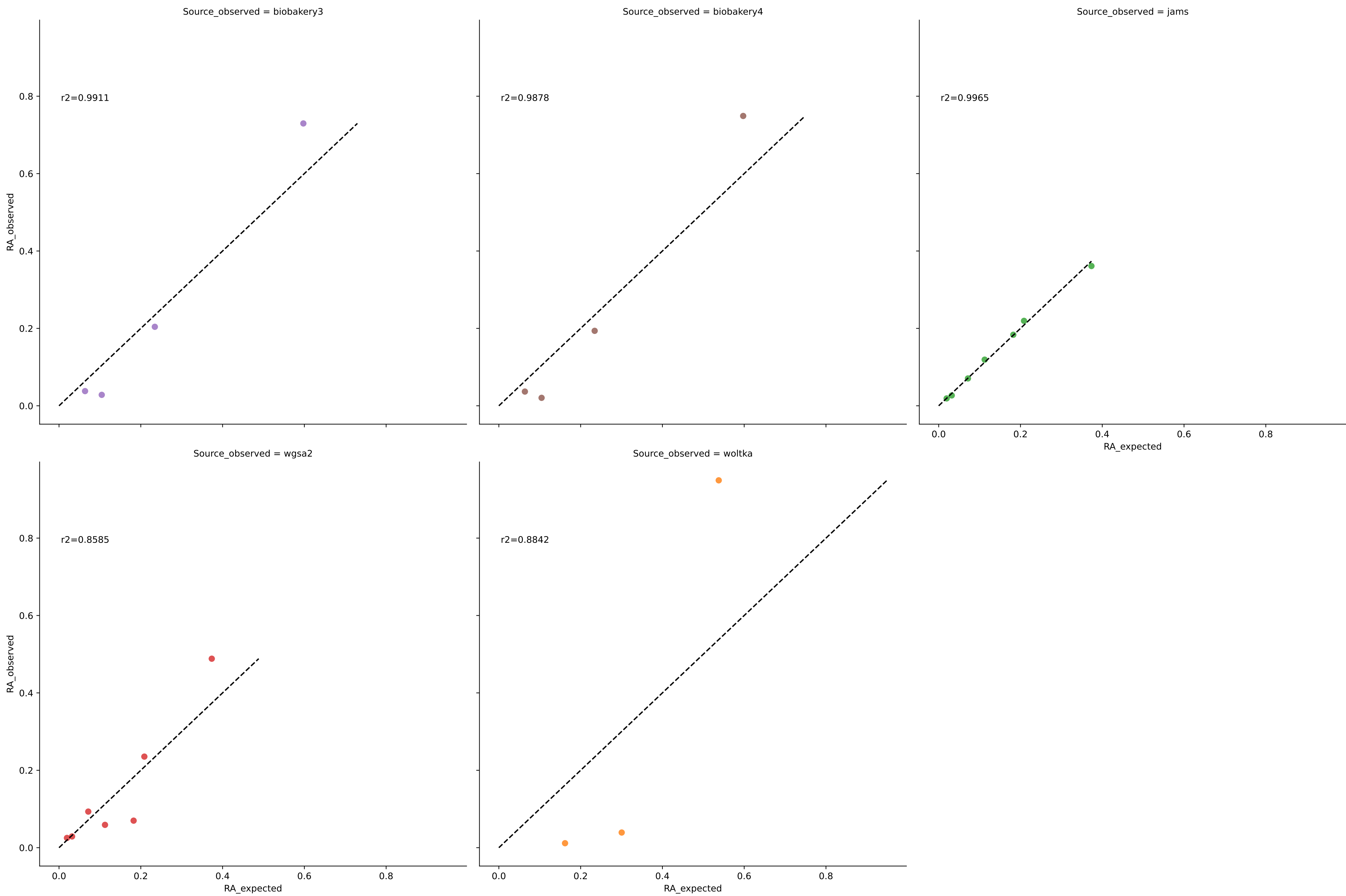
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487937	16	0.6864	0.0042	1.8398	0.8307	0.0055	100.0000	0.0000
SRR11487938	16	0.6953	0.0042	1.8839	0.8305	0.0056	100.0000	0.0000
SRR11487939	16	0.7025	0.0042	1.8862	0.8302	0.0056	100.0000	0.0000
SRR11487940	16	0.6998	0.0042	1.8456	0.8314	0.0056	100.0000	0.0000
SRR11487941	16	0.7008	0.0042	1.8543	0.8316	0.0056	100.0000	0.0000
Average	16	0.6969	0.0042	1.8620	0.8309	0.0056	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using wol in Experiment Amos mixed with filter 0.001

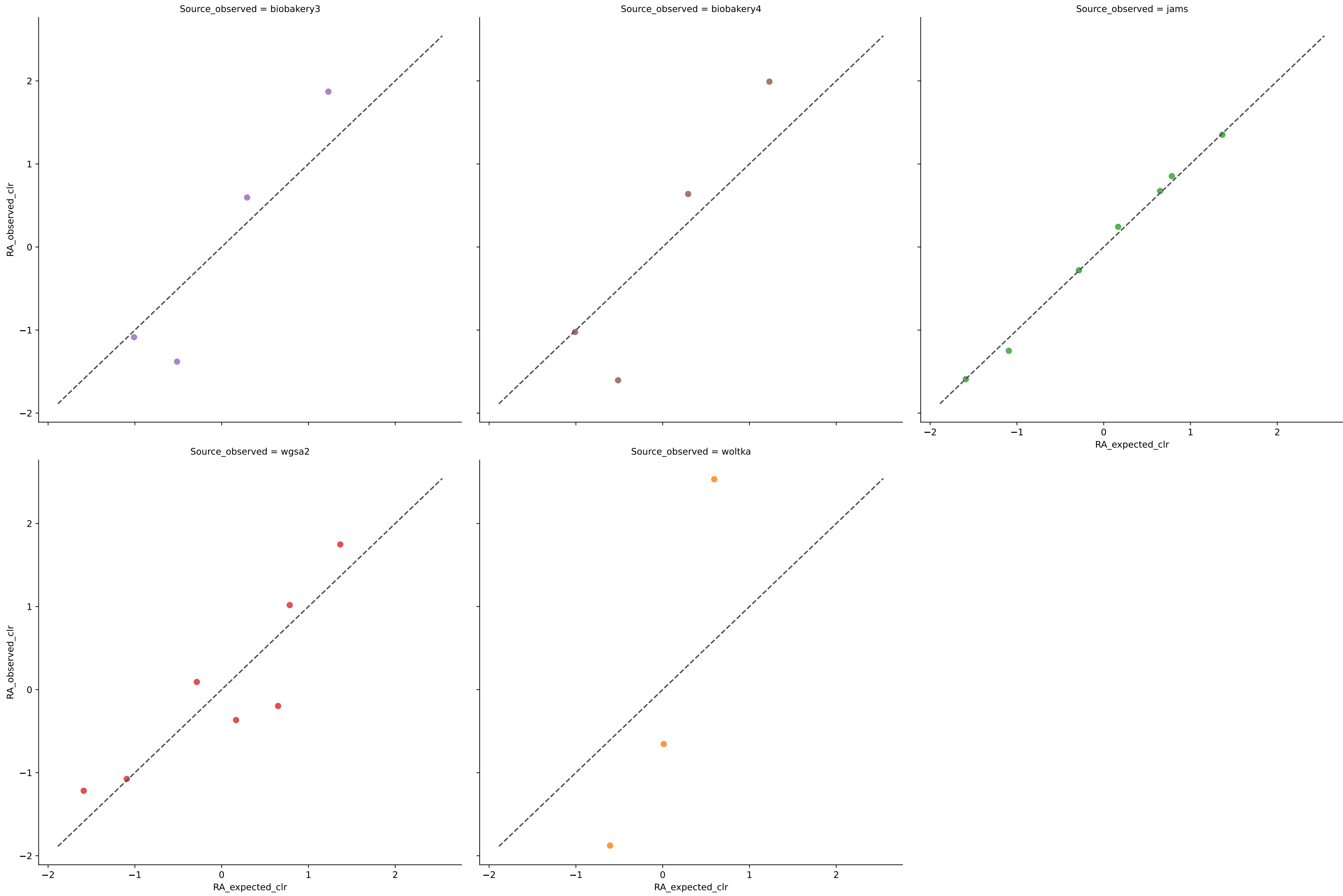


	Diversity	R ²	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487937	18	0.0234	0.0058	3.2948	0.7376	0.0085	100.0000	0.0000
SRR11487938	18	0.0204	0.0058	3.3182	0.7373	0.0085	100.0000	0.0000
SRR11487939	18	0.0221	0.0058	3.3334	0.7377	0.0084	100.0000	0.0000
SRR11487940	18	0.0196	0.0058	3.2964	0.7397	0.0084	100.0000	0.0000
SRR11487941	18	0.0193	0.0058	3.2935	0.7407	0.0084	100.0000	0.0000
Average	18	0.0210	0.0058	3.3073	0.7386	0.0085	100.0000	0.0000

Bivariate Linear Regression for Sample S1 in Experiment bmock12 (Genus at filter threshold 0.01)

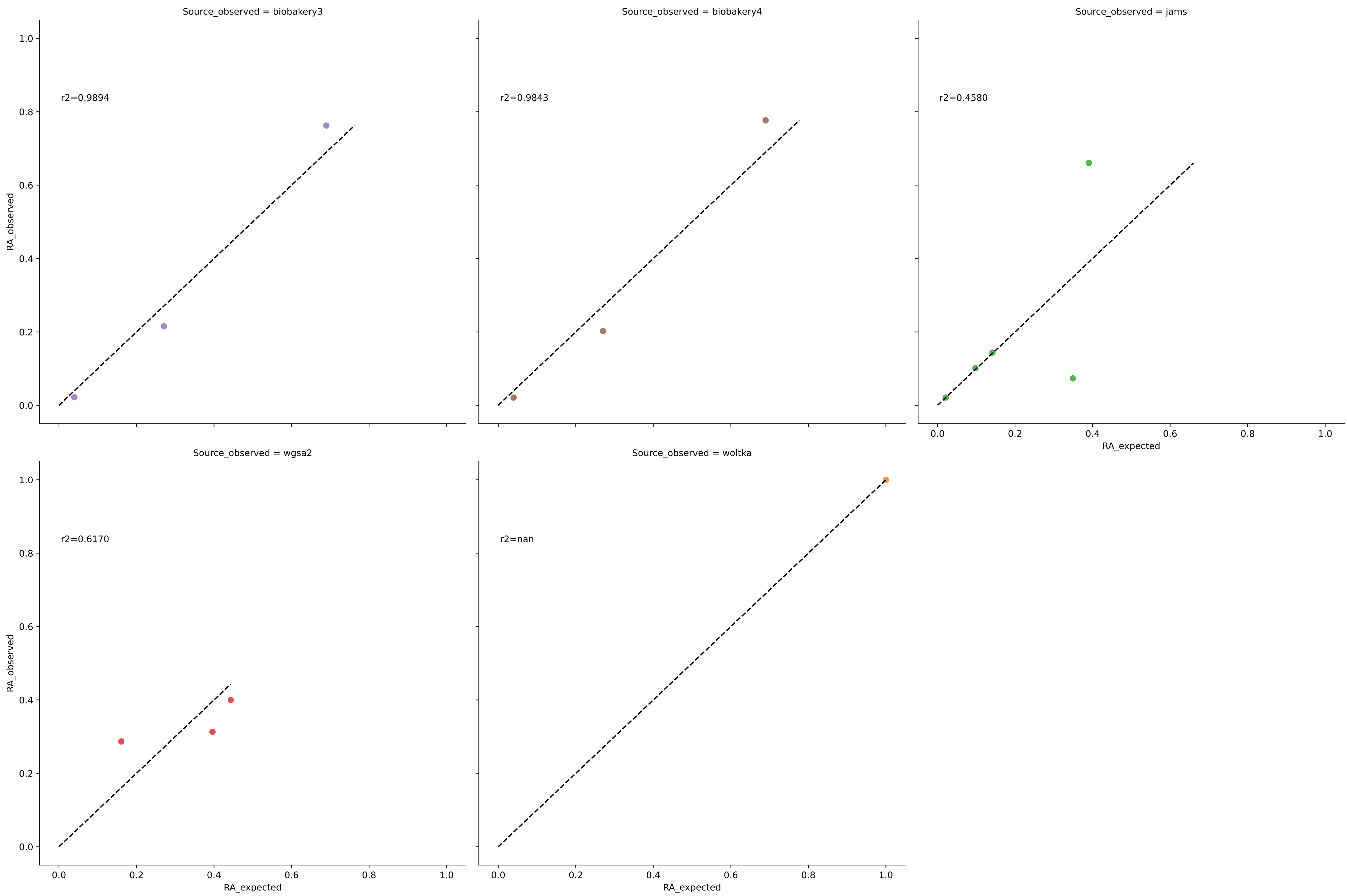


Bivariate Linear Regression for Sample S1 in Experiment bmock12 (Genus at filter threshold 0.01)

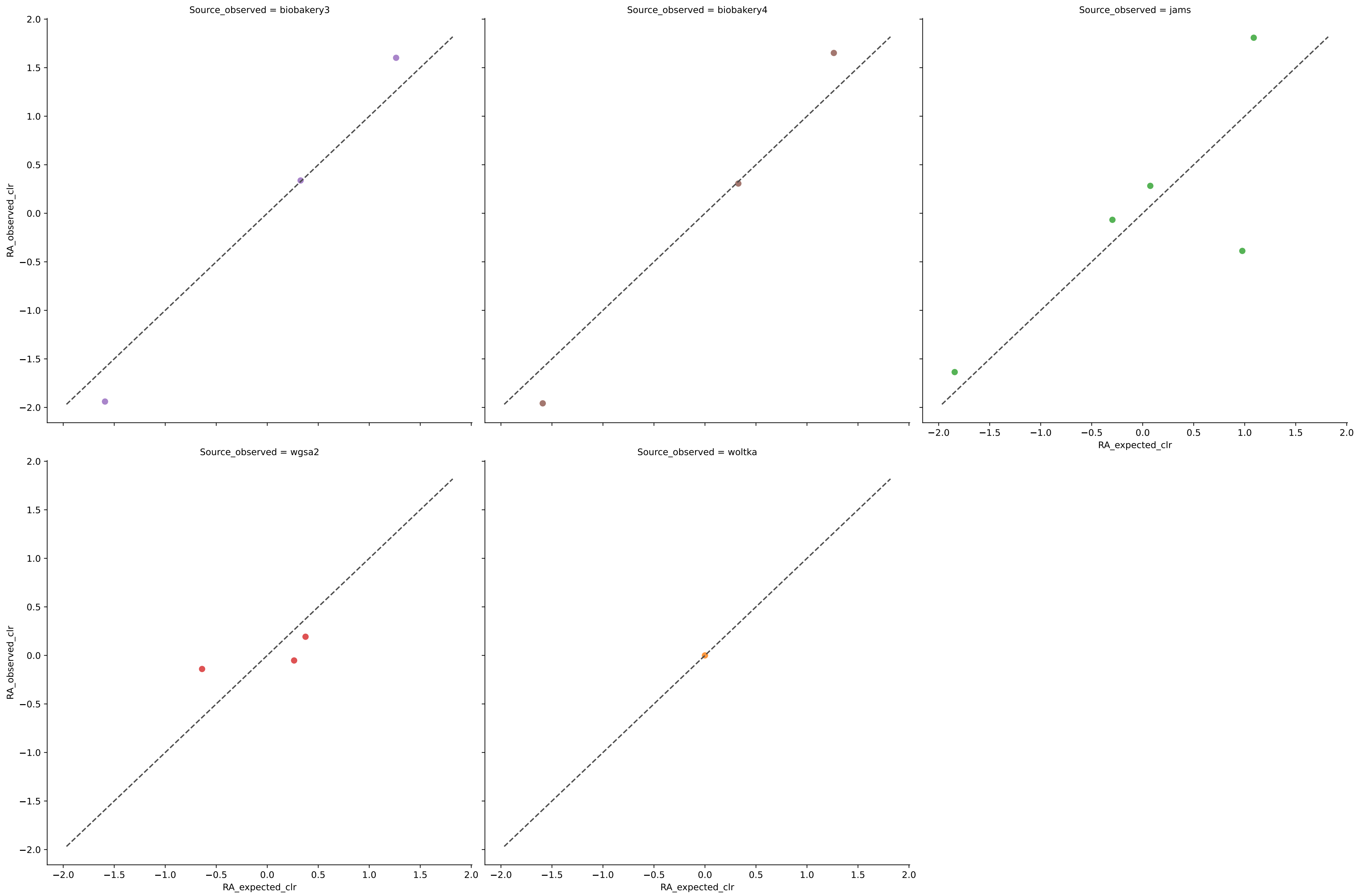


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
biobakery3	4	0.9911	0.0661	1.1219	0.8679	0.0788	100.0000	0.0000
biobakery4	4	0.9878	0.0757	1.3740	0.8486	0.0899	100.0000	0.0000
jams	7	0.9965	0.0053	0.1890	0.9813	0.0070	100.0000	0.0000
wgsa2	7	0.8585	0.0483	1.2171	0.8311	0.0654	100.0000	0.0000
woltka	3	0.8842	0.2744	2.4122	0.5884	0.2945	100.0000	0.0000

Bivariate Linear Regression for Sample S1 in Experiment bmock12 (Species at filter threshold 0.01)

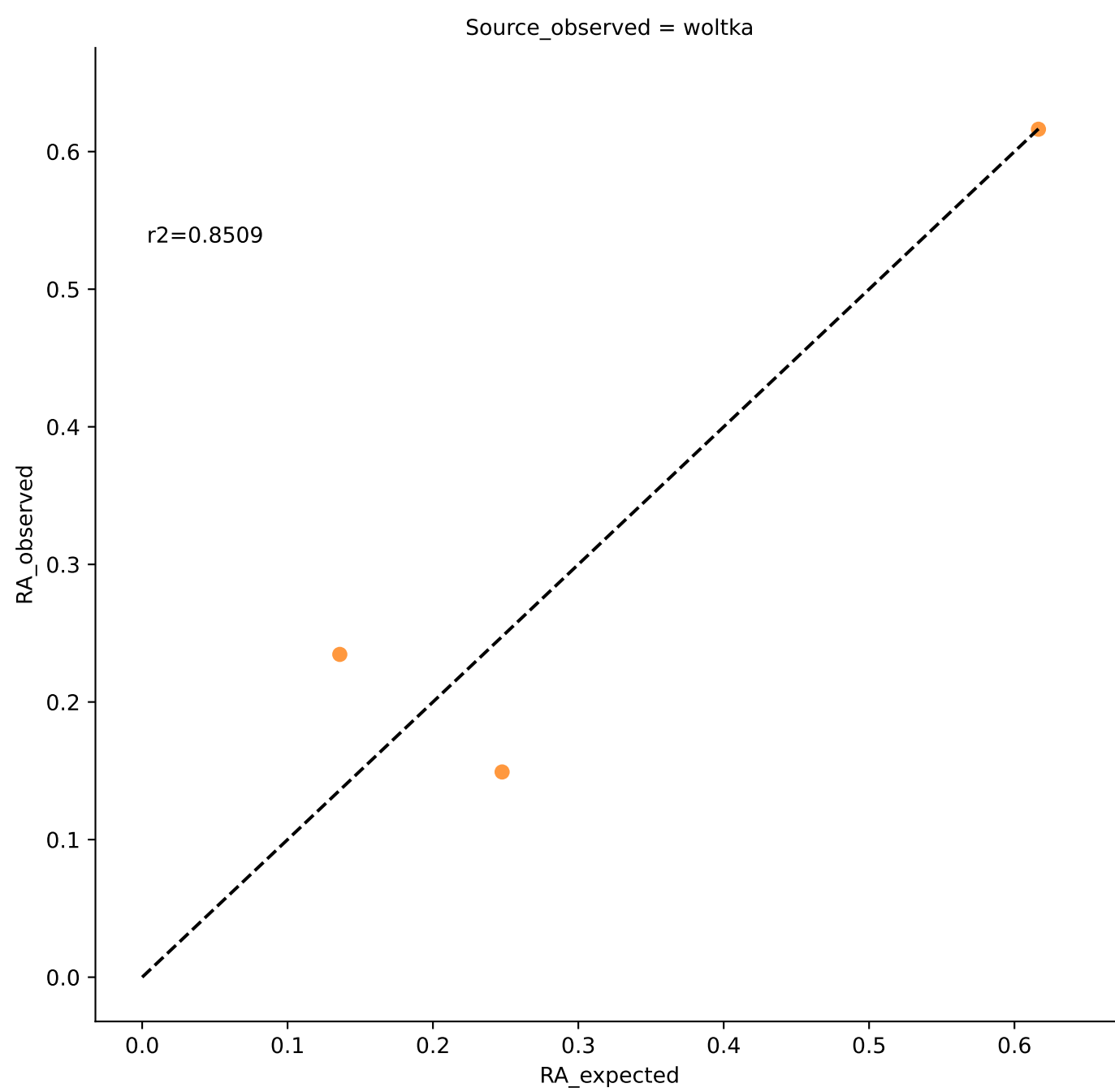
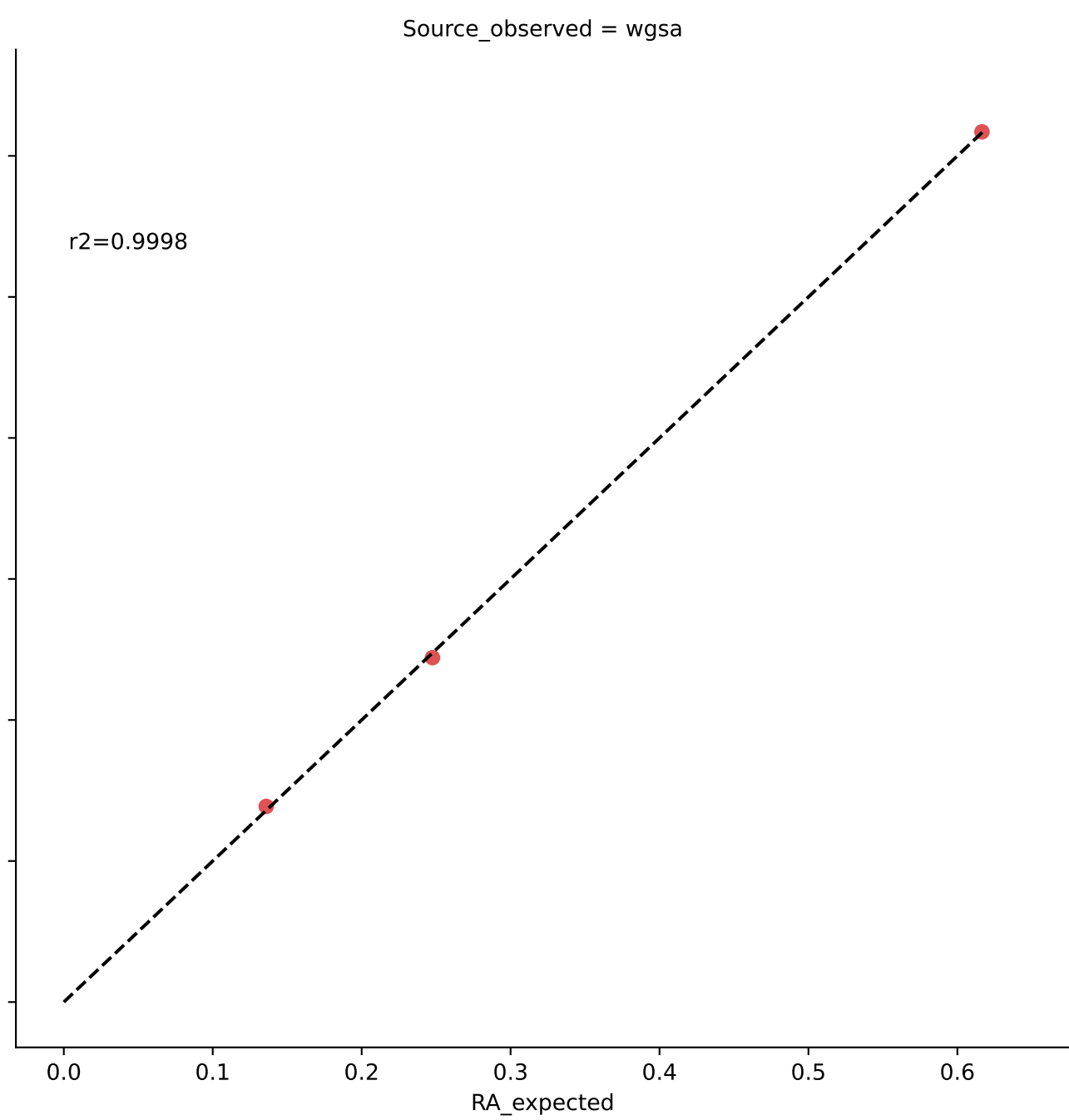
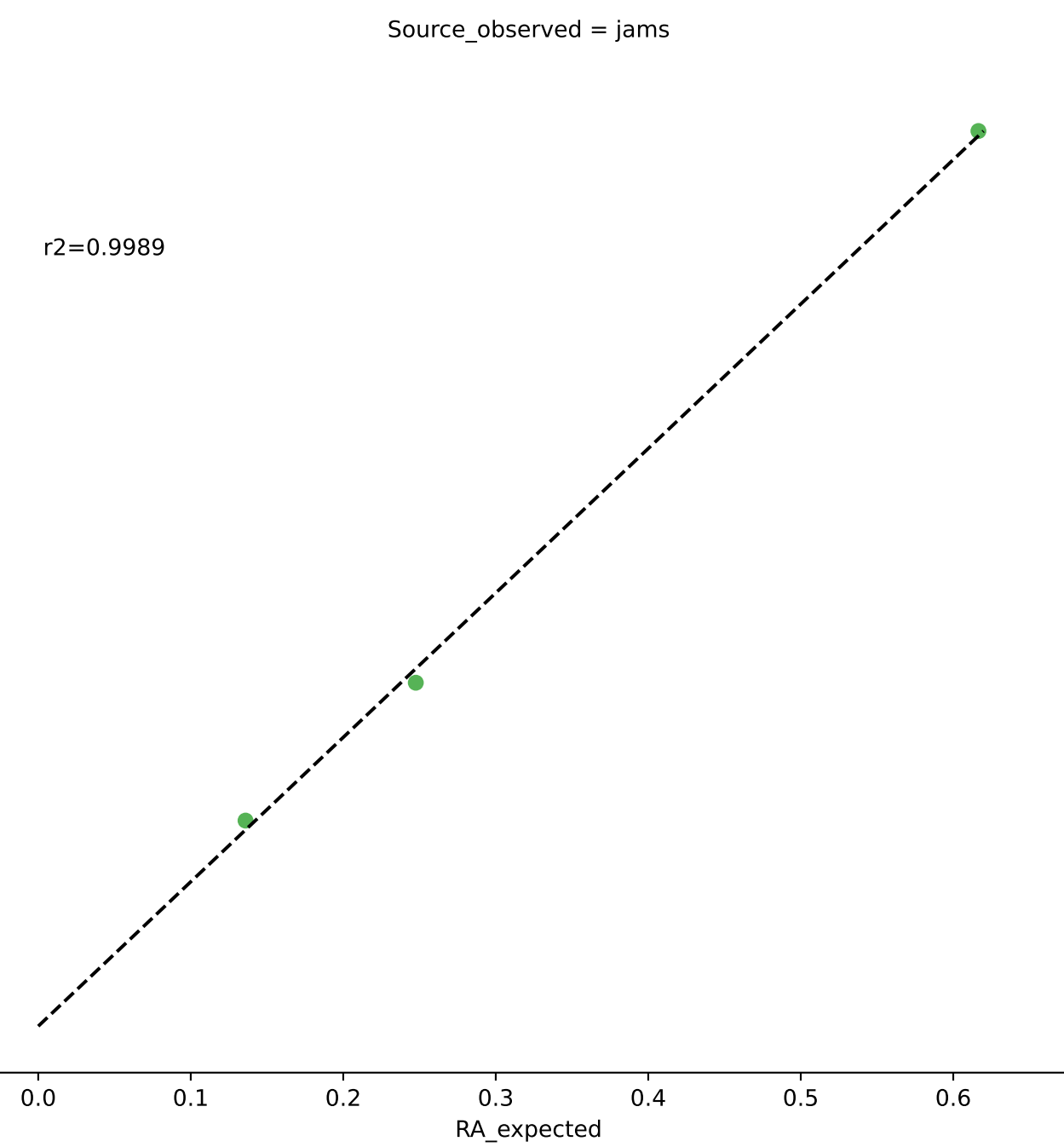
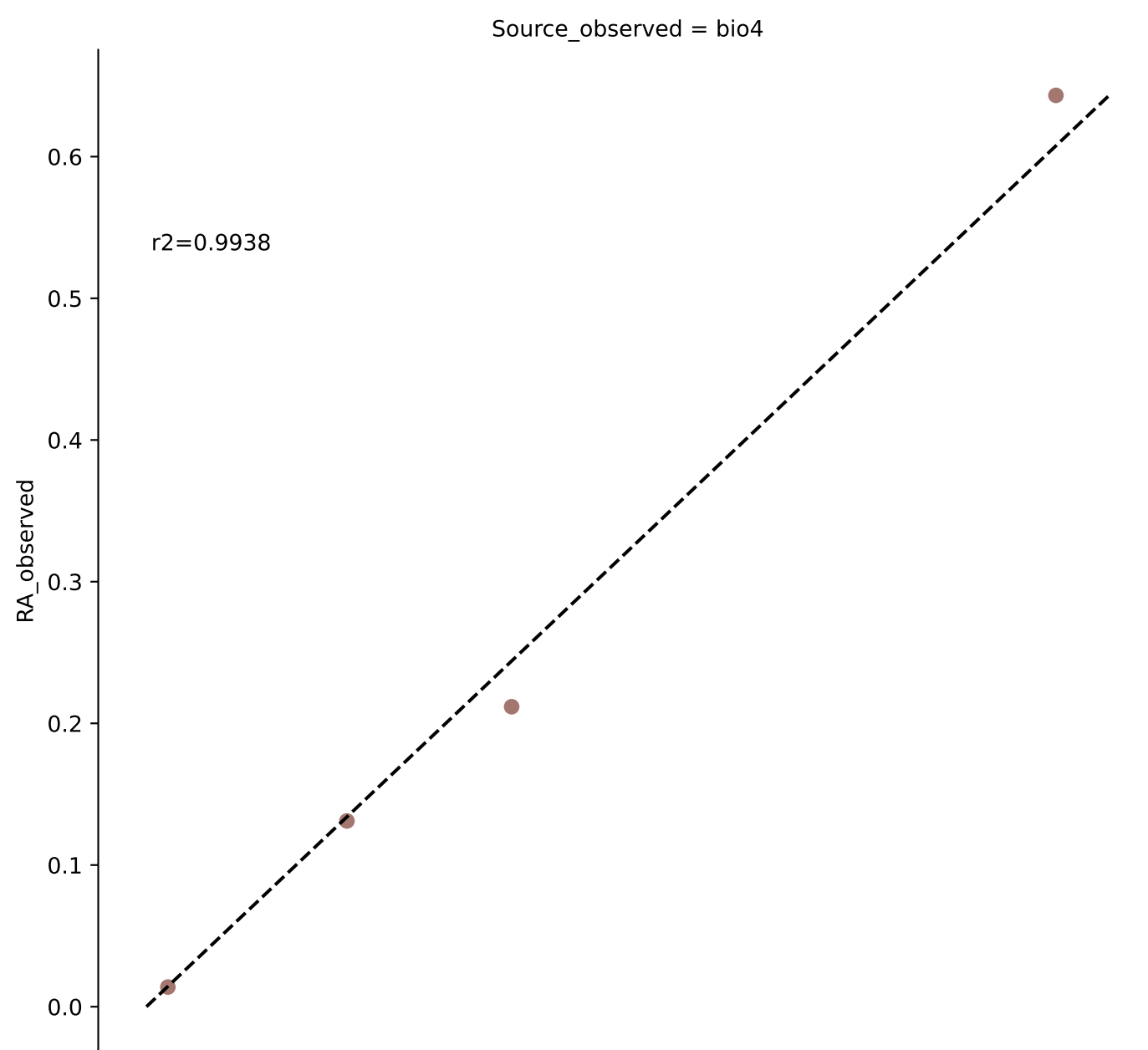


Bivariate Linear Regression for Sample S1 in Experiment bmock12 (Species at filter threshold 0.01)

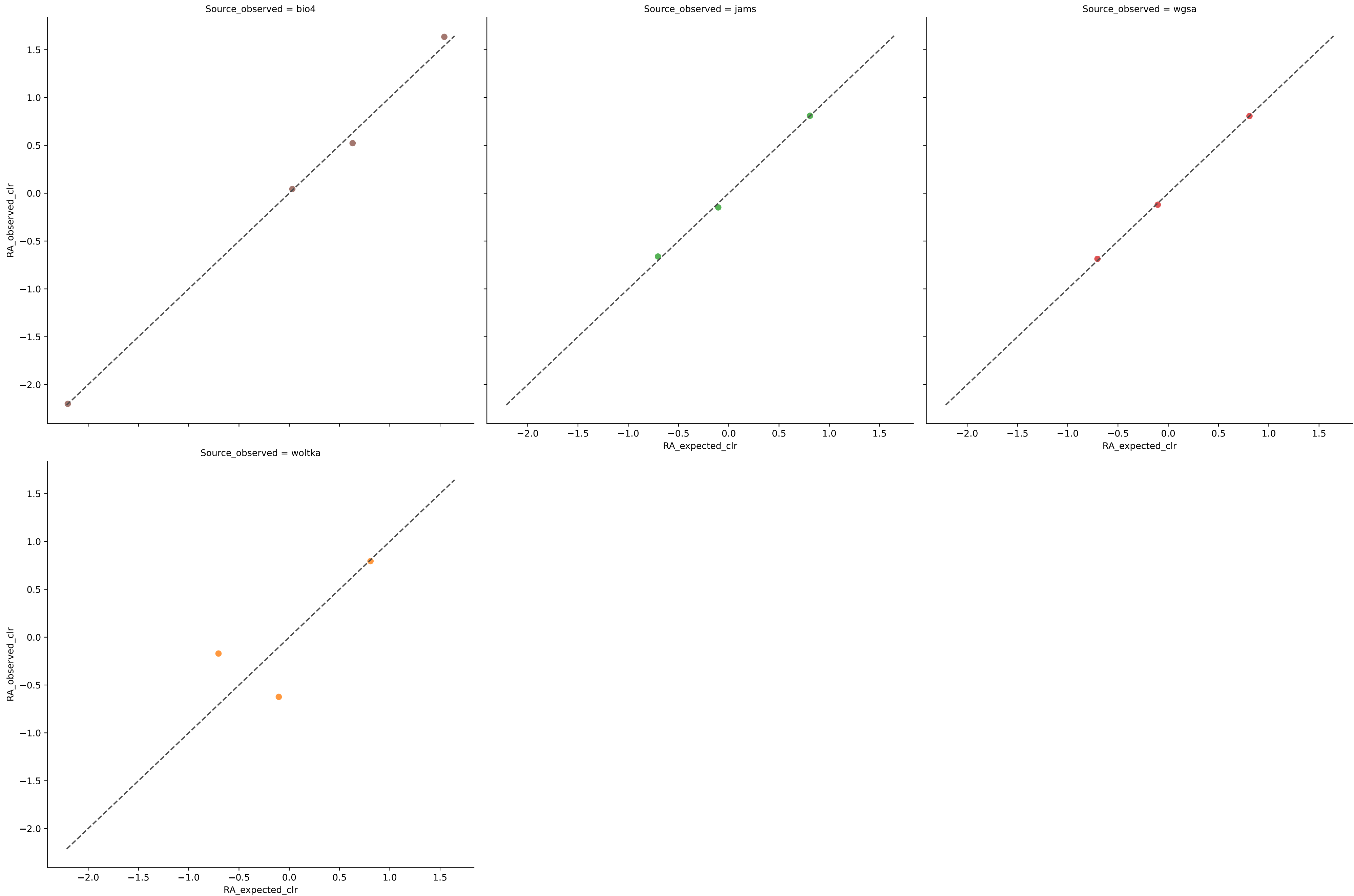


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
biobakery3	3	0.9894	0.0485	0.4862	0.9273	0.0536	100.0000	0.0000
biobakery4	3	0.9843	0.0579	0.5350	0.9131	0.0647	100.0000	0.0000
jams	5	0.4580	0.1103	1.5869	0.7243	0.1726	100.0000	0.0000
wgsa2	3	0.6170	0.0841	0.6180	0.8738	0.0907	100.0000	0.0000
woltka	1	nan	0.0000	0.0000	1.0000	0.0000	100.0000	0.0000

Bivariate Linear Regression for Sample S1 in Experiment camisimGI (Genus at filter threshold 0.01)

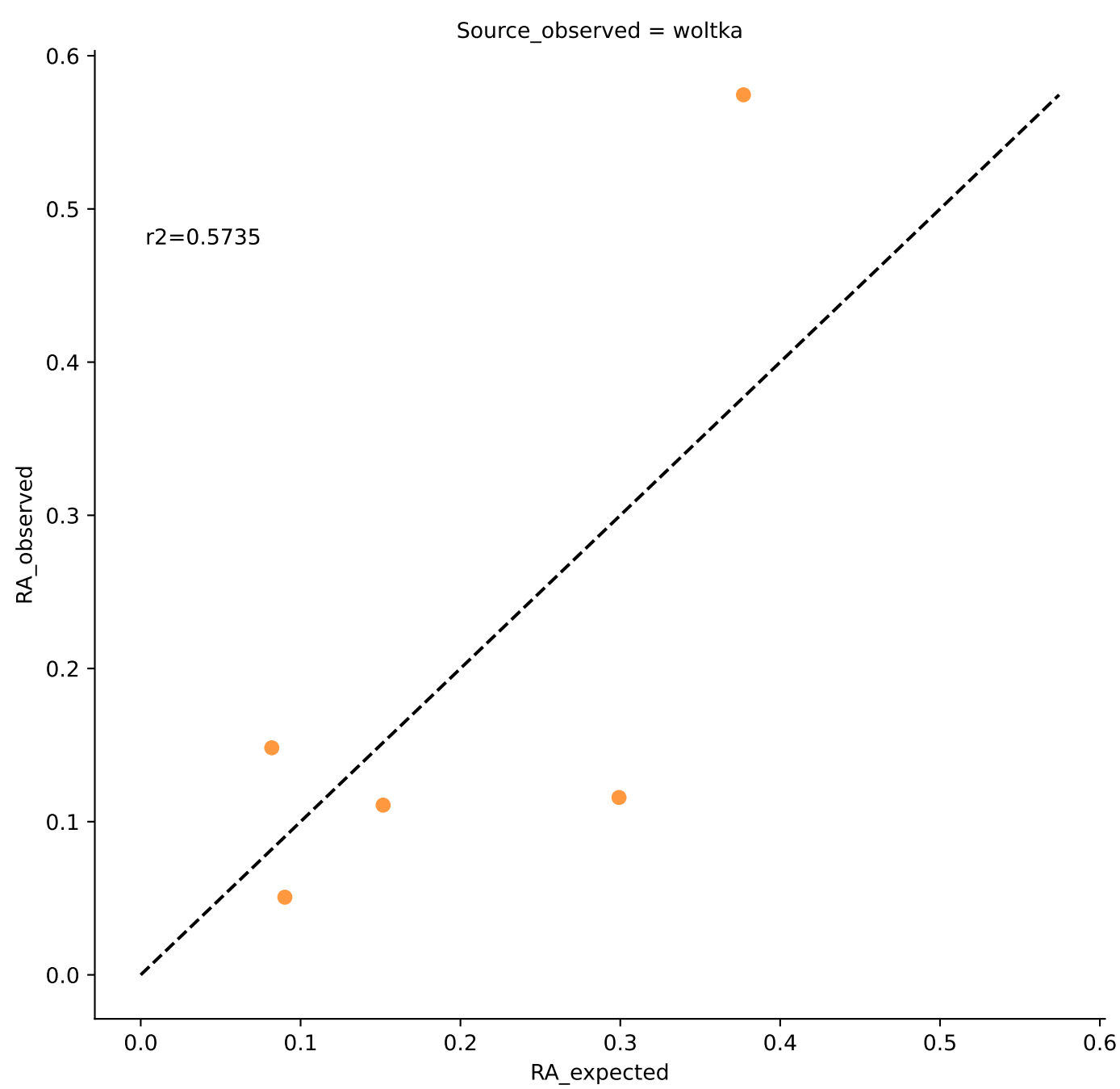
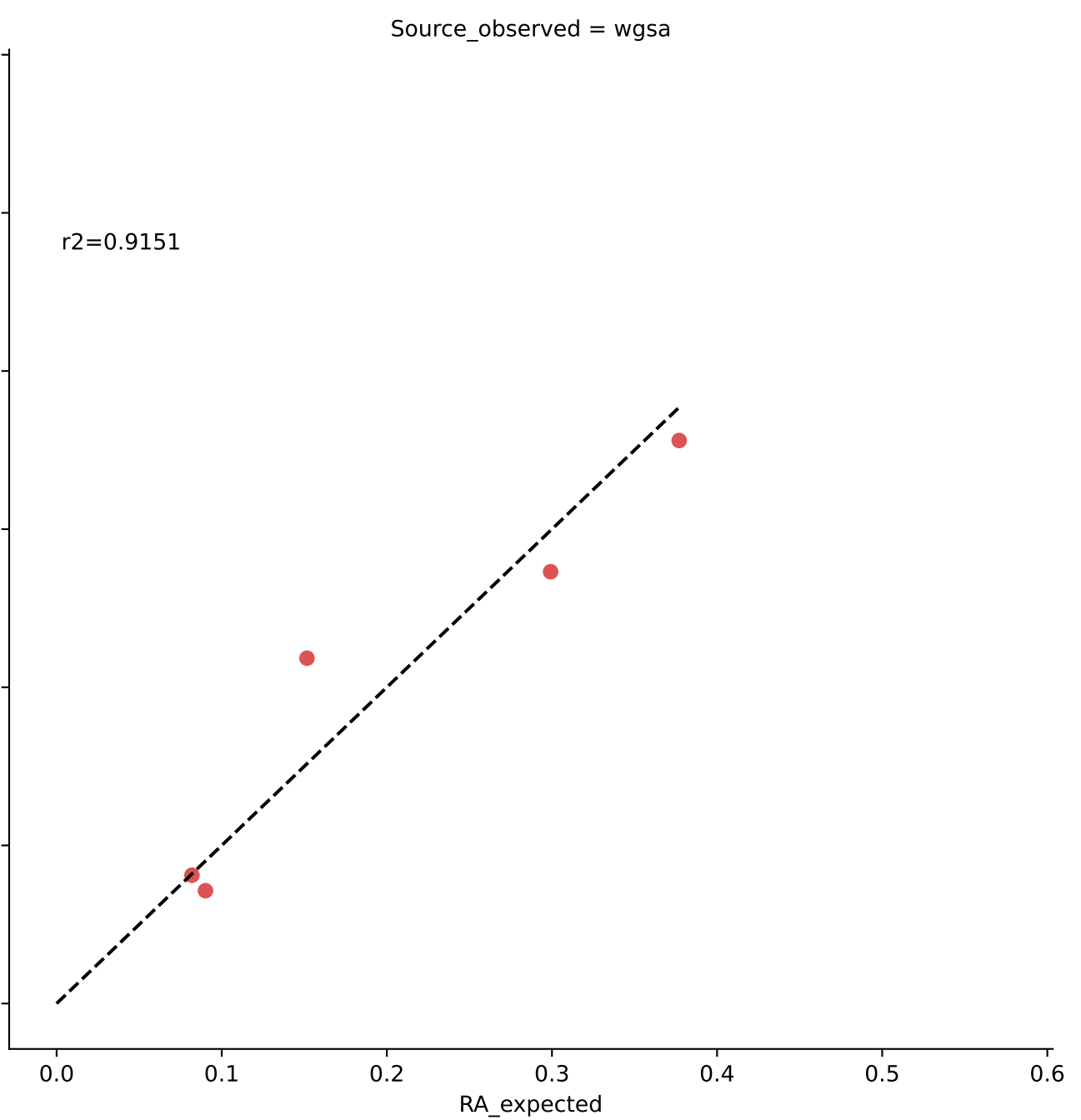
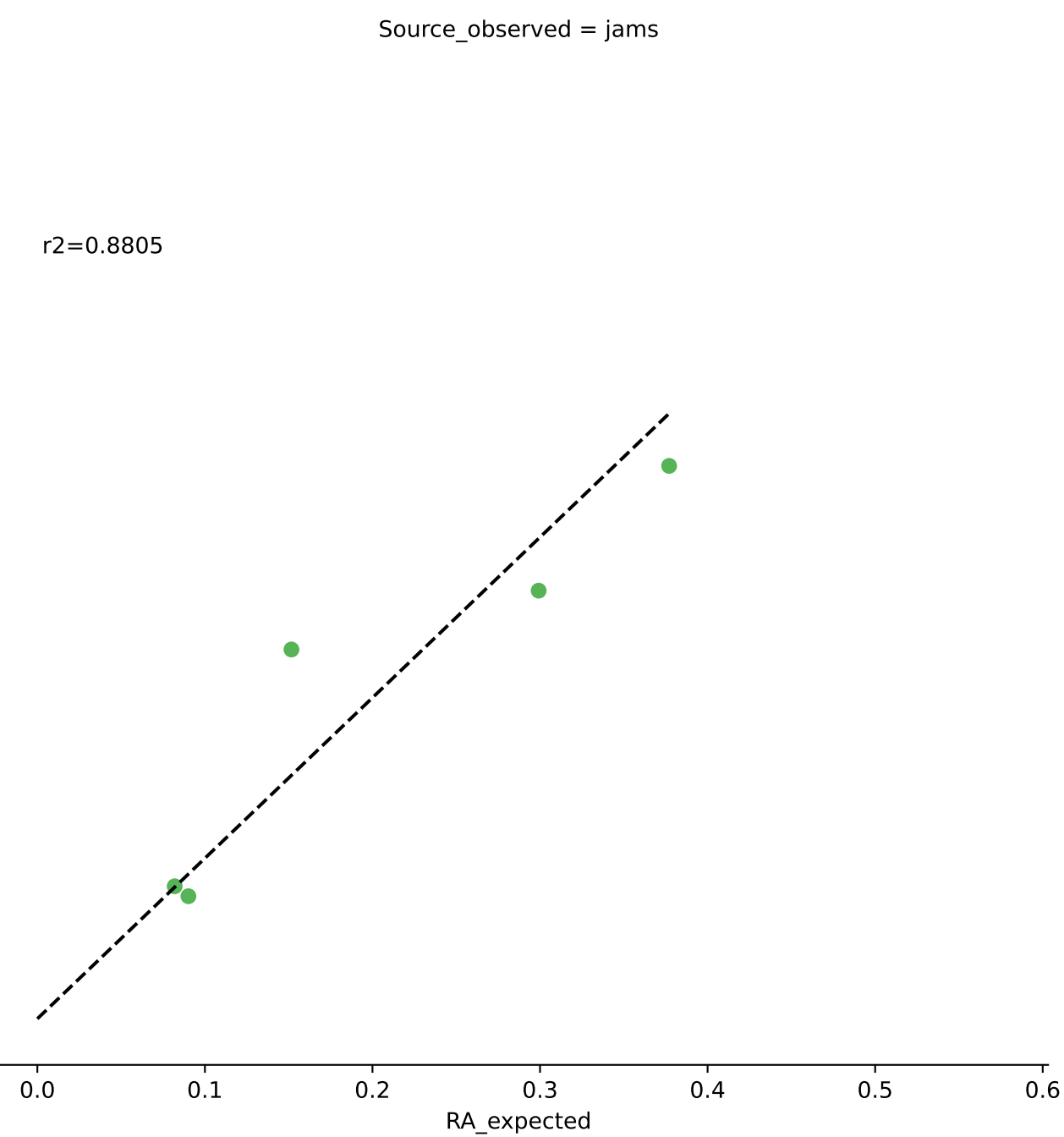
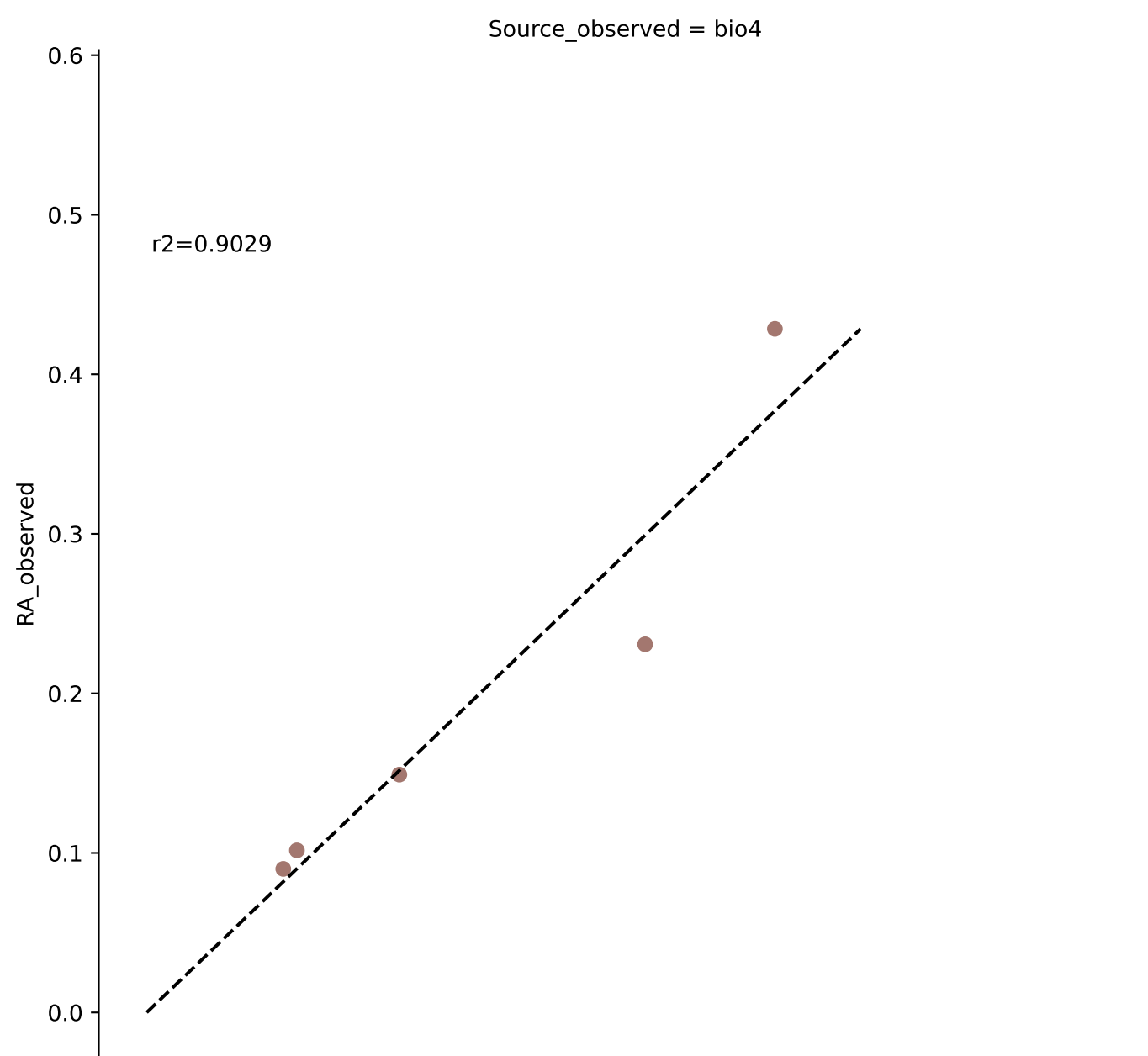


Bivariate Linear Regression for Sample S1 in Experiment camisimGI (Genus at filter threshold 0.01)

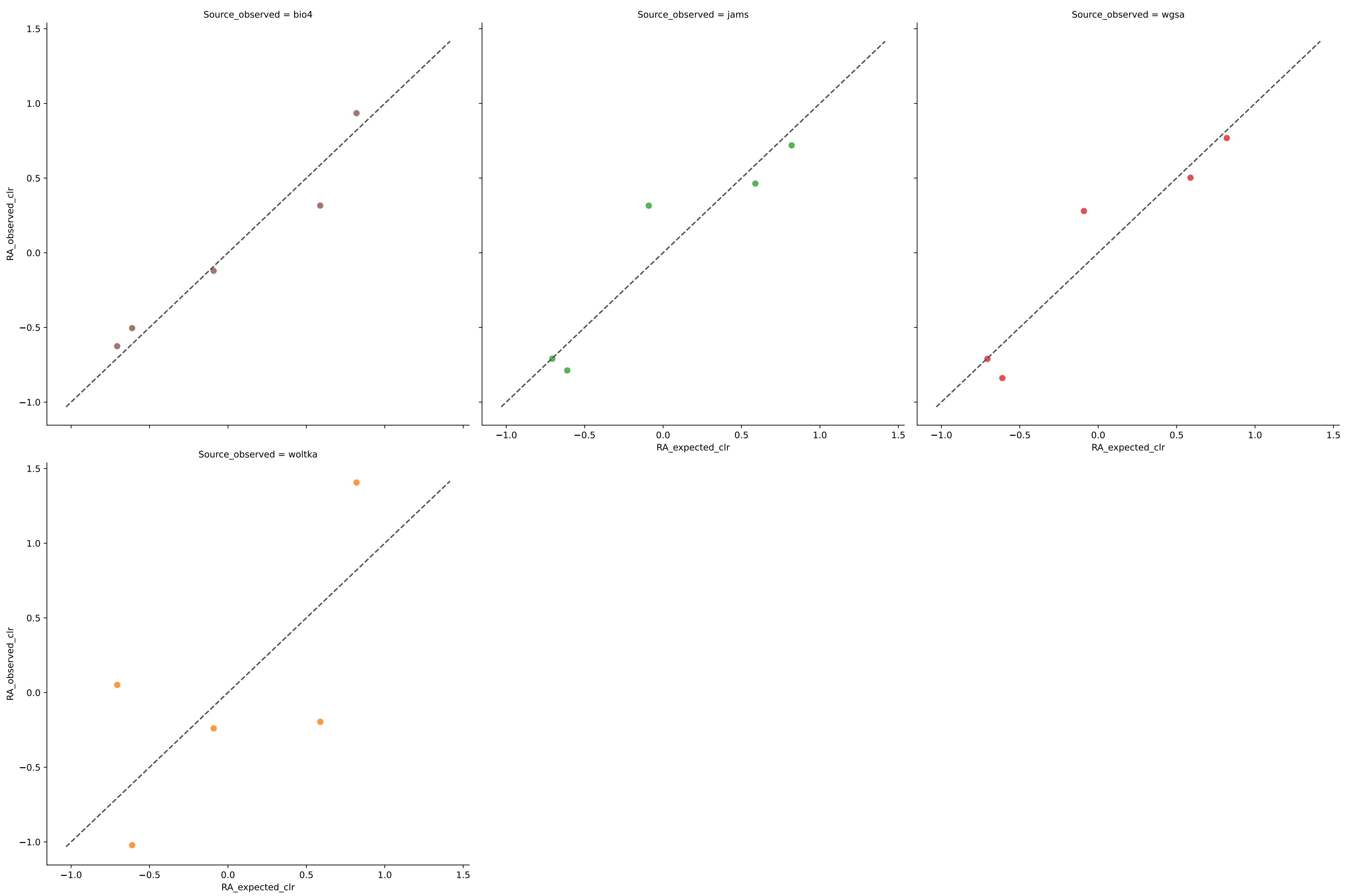


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	4	0.9938	0.0178	0.1416	0.9644	0.0241	100.0000	0.0000
jams	3	0.9989	0.0065	0.0612	0.9902	0.0070	100.0000	0.0000
wgsa	3	0.9998	0.0022	0.0242	0.9966	0.0025	100.0000	0.0000
woltka	3	0.8509	0.0658	0.7444	0.9013	0.0805	100.0000	0.0000

Bivariate Linear Regression for Sample S2 in Experiment camisimGI (Genus at filter threshold 0.01)

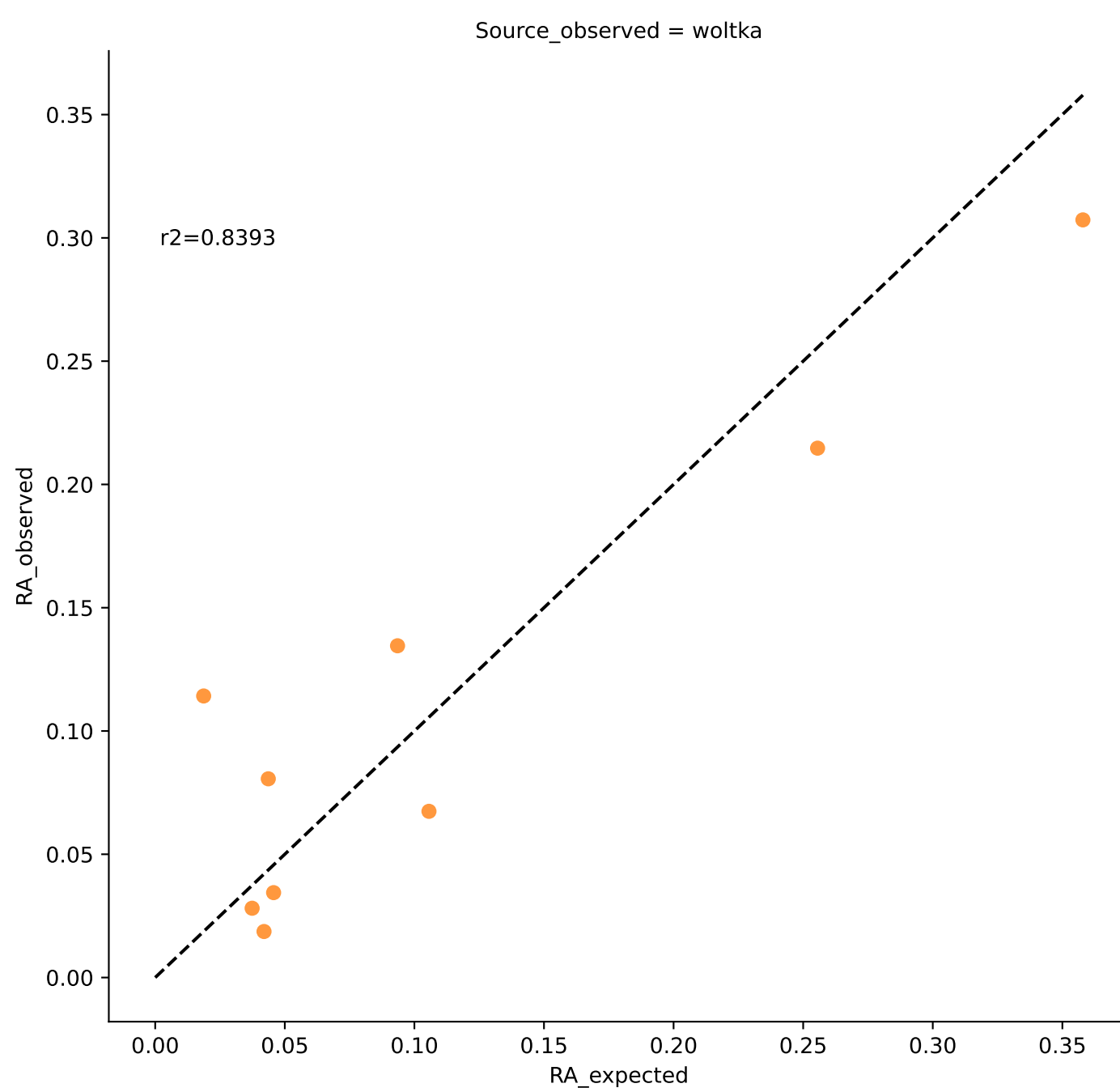
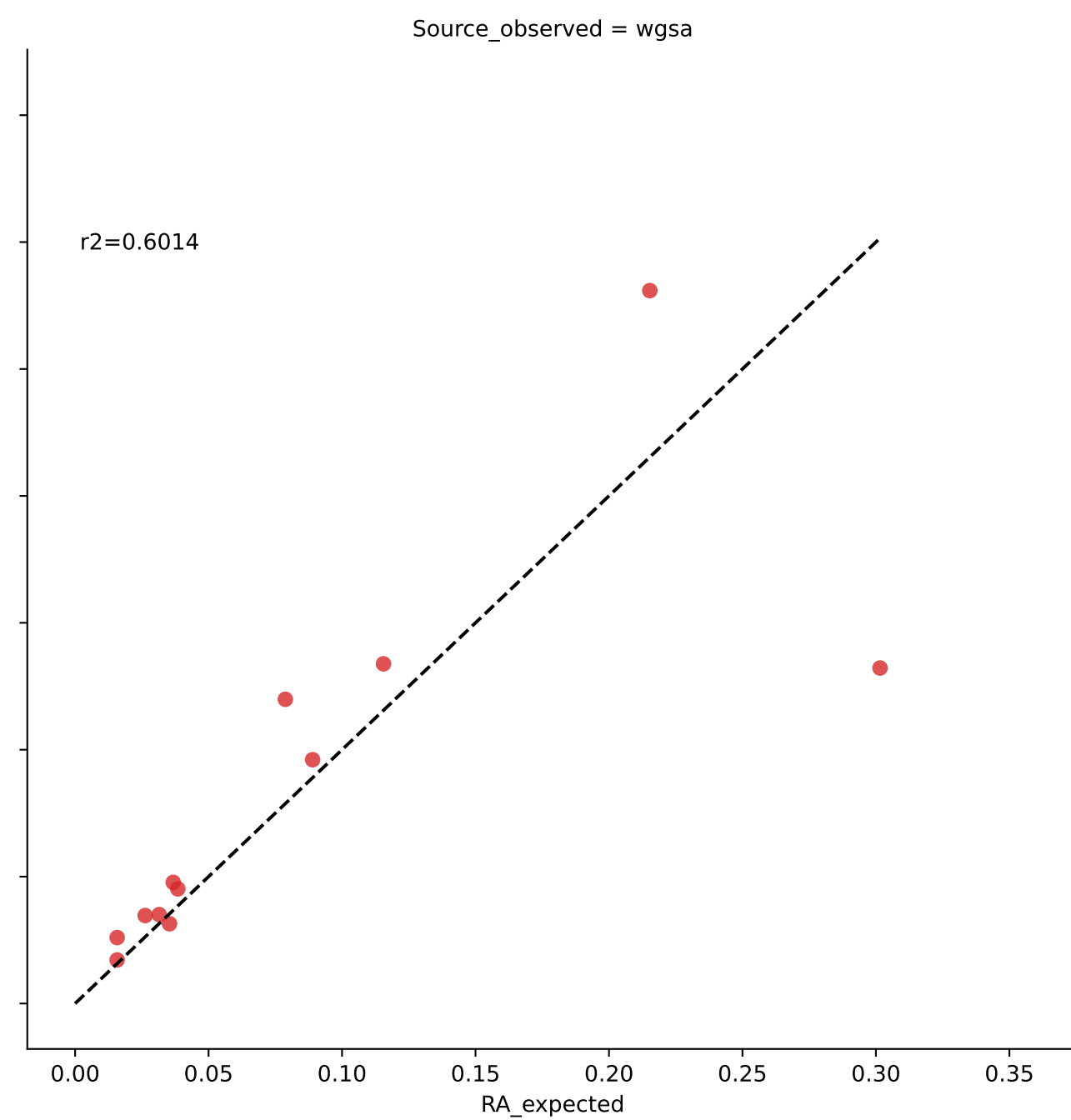
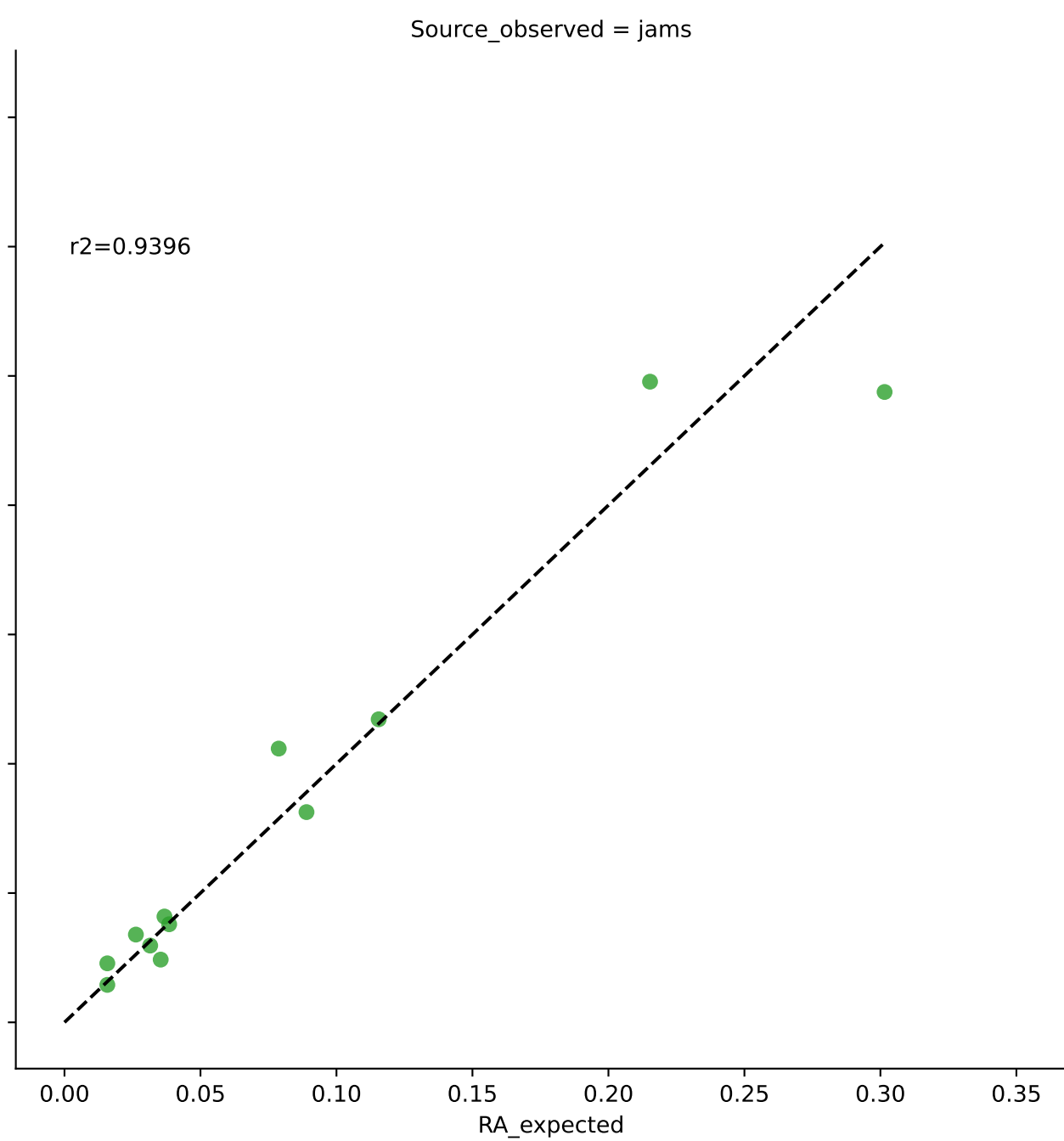
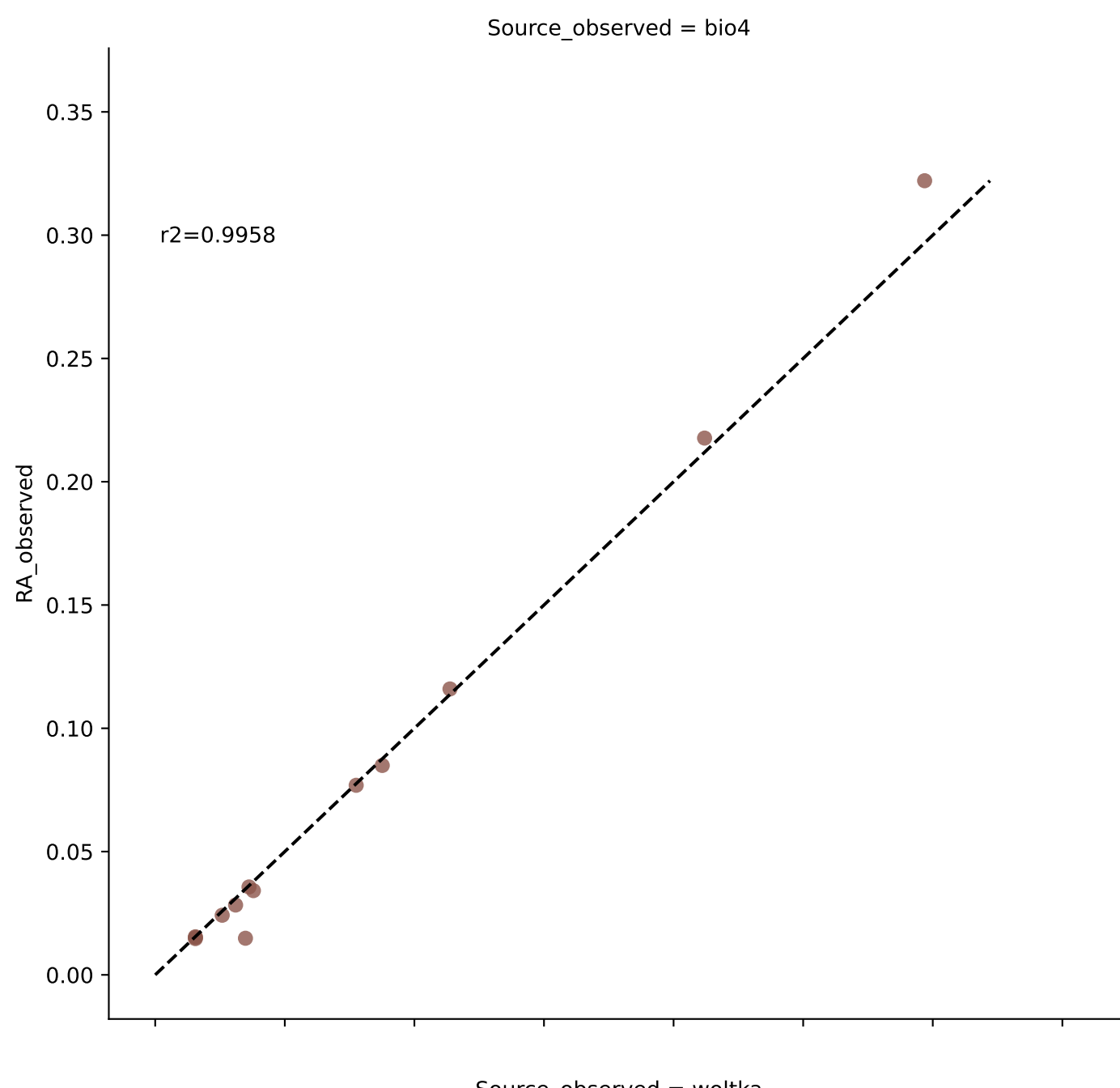


Bivariate Linear Regression for Sample S2 in Experiment camisimGI (Genus at filter threshold 0.01)

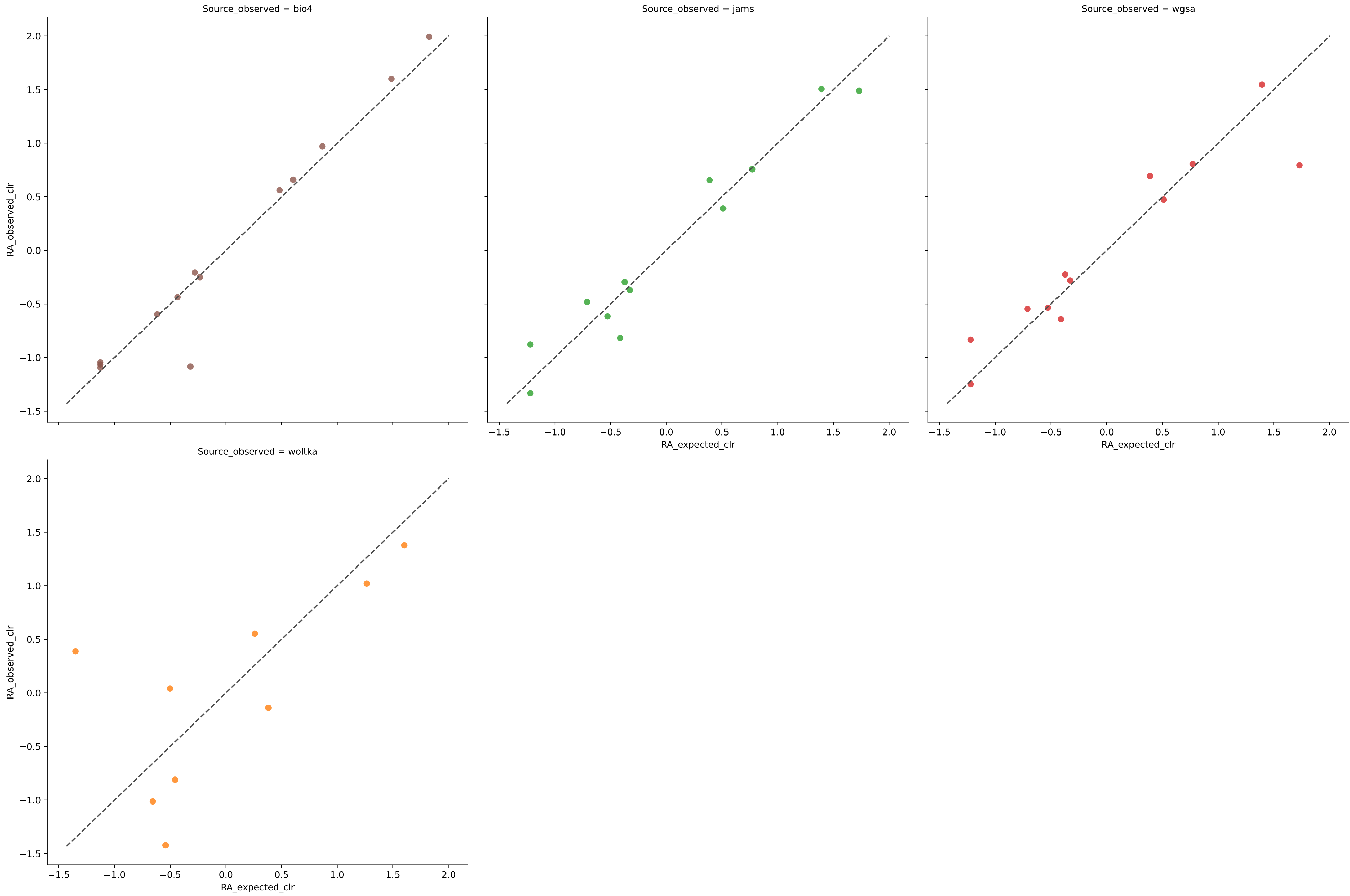


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	5	0.9029	0.0284	0.3259	0.9291	0.0388	100.0000	0.0000
jams	5	0.8805	0.0316	0.4715	0.9210	0.0411	100.0000	0.0000
wgsa	5	0.9151	0.0267	0.4465	0.9332	0.0345	100.0000	0.0000
woltka	5	0.5735	0.1055	1.3129	0.7363	0.1267	100.0000	0.0000

Bivariate Linear Regression for Sample S1 in Experiment camisimGI (Species at filter threshold 0.01)

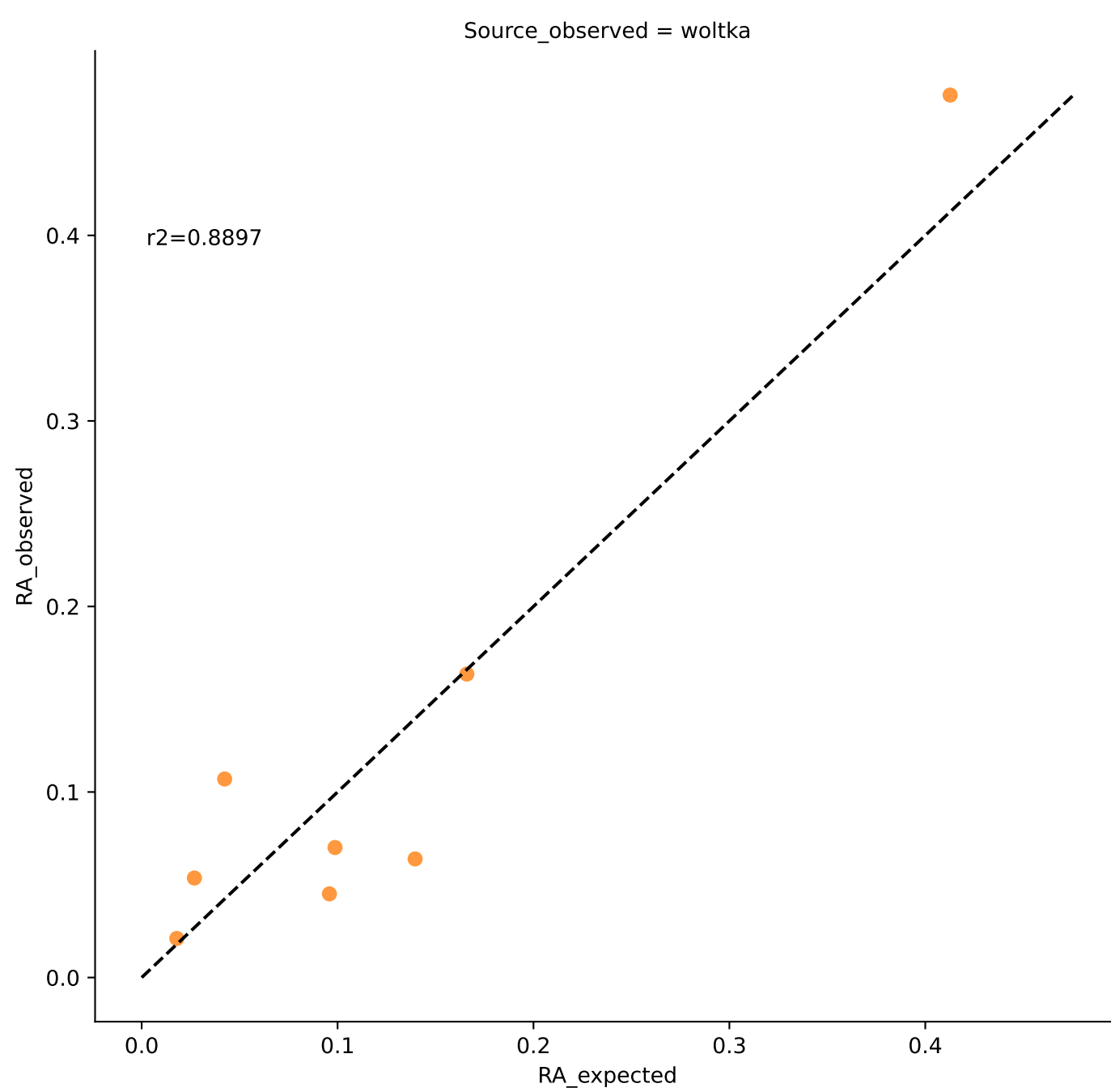
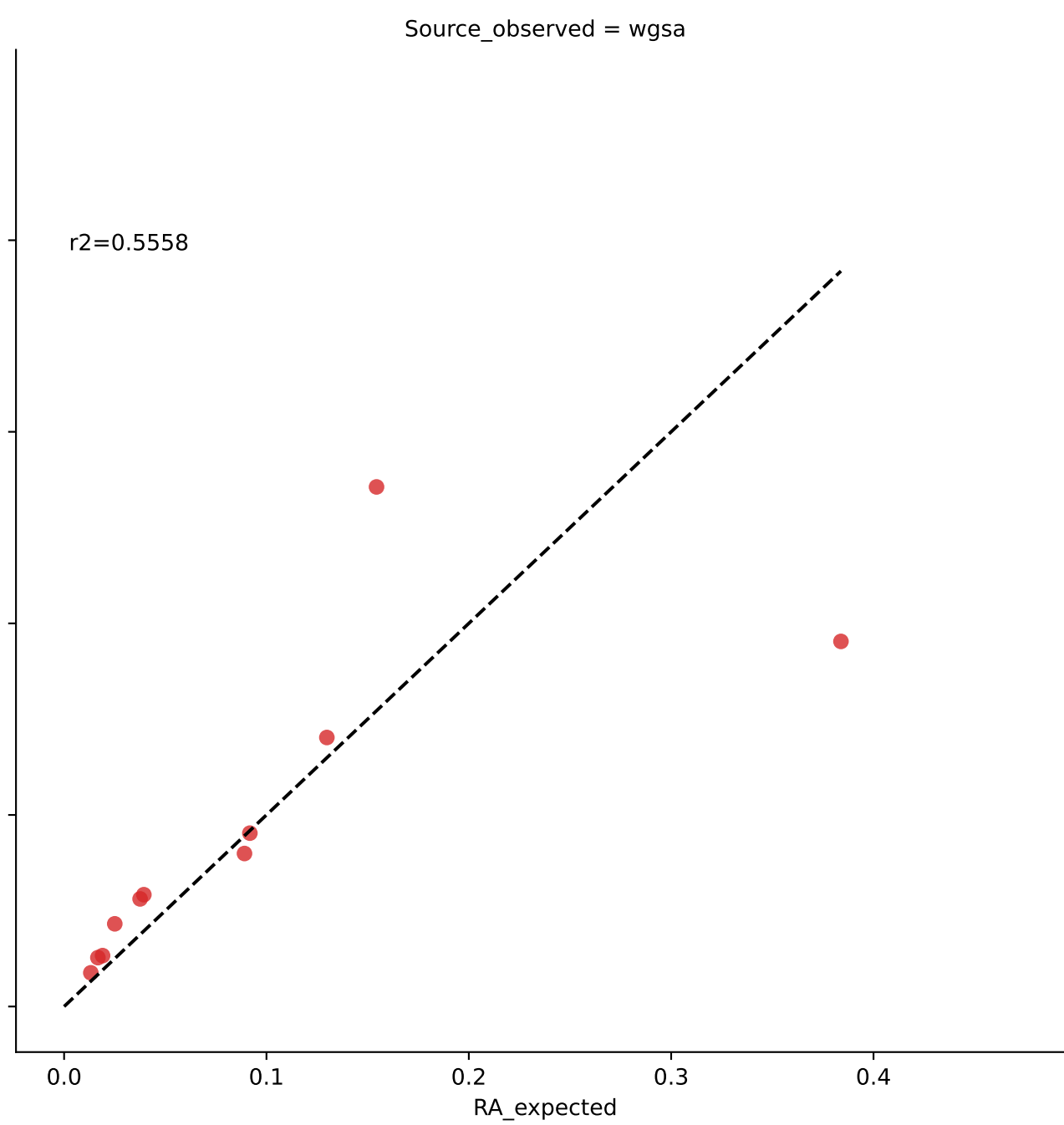
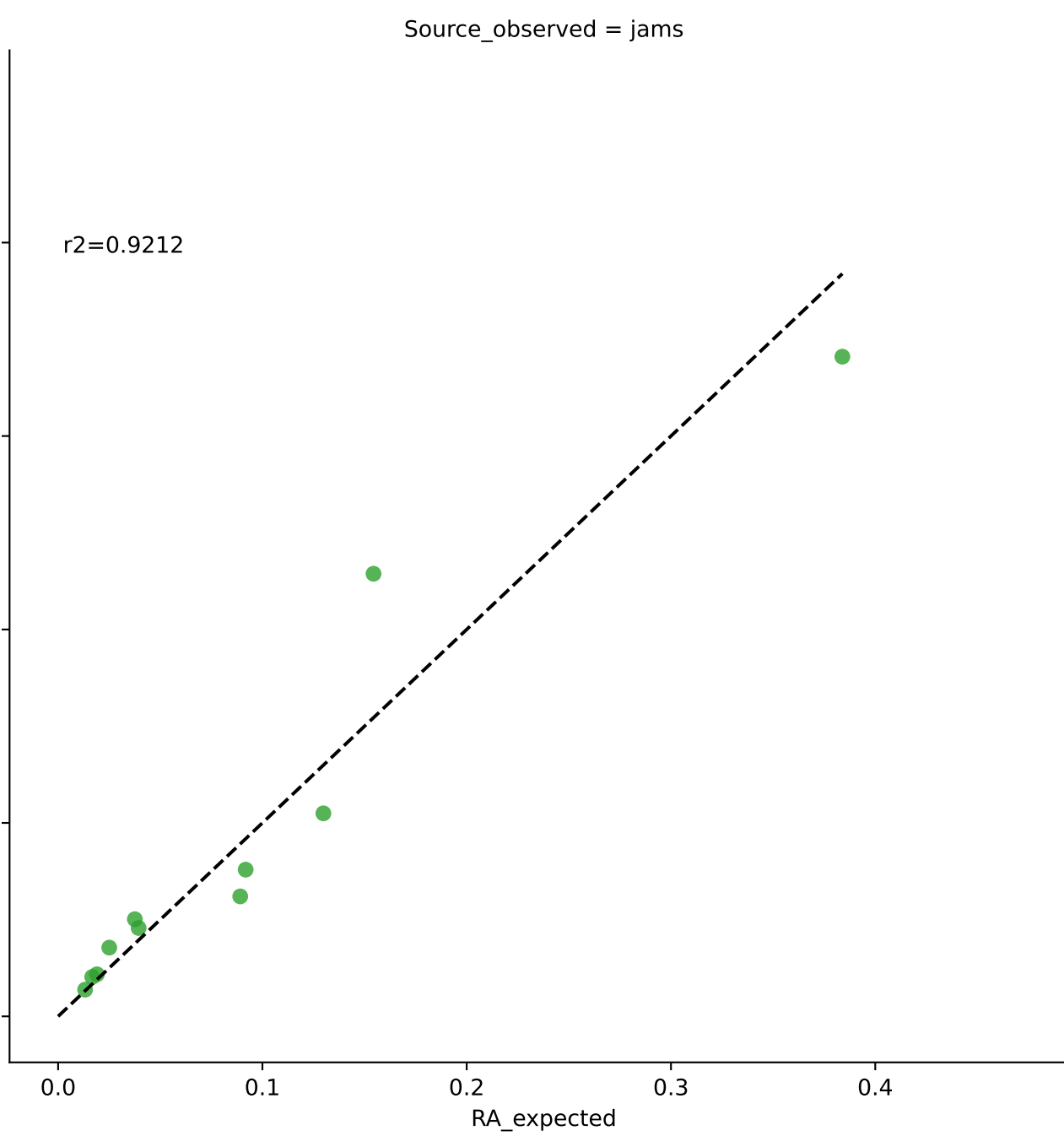
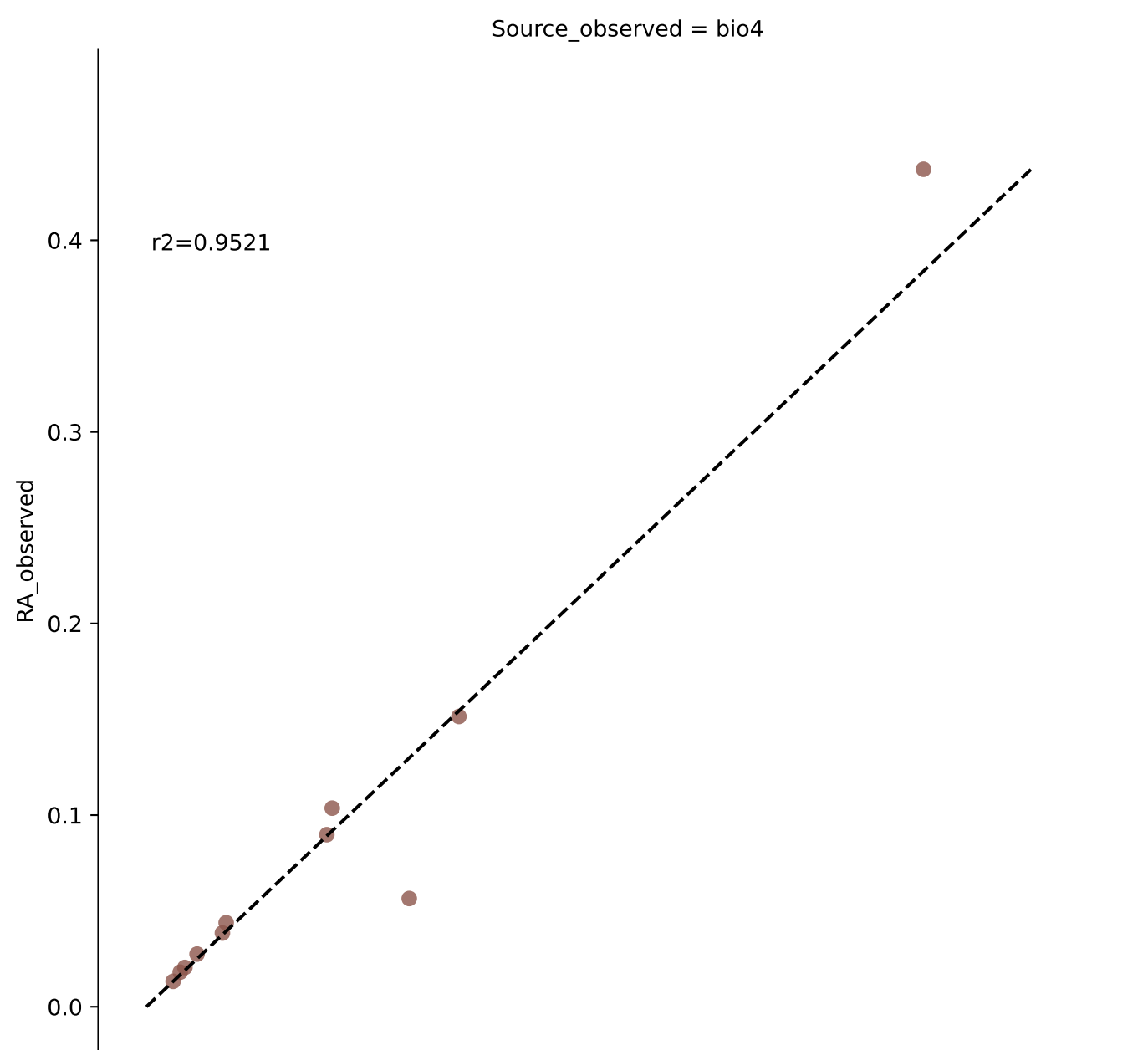


Bivariate Linear Regression for Sample S1 in Experiment camisimGI (Species at filter threshold 0.01)

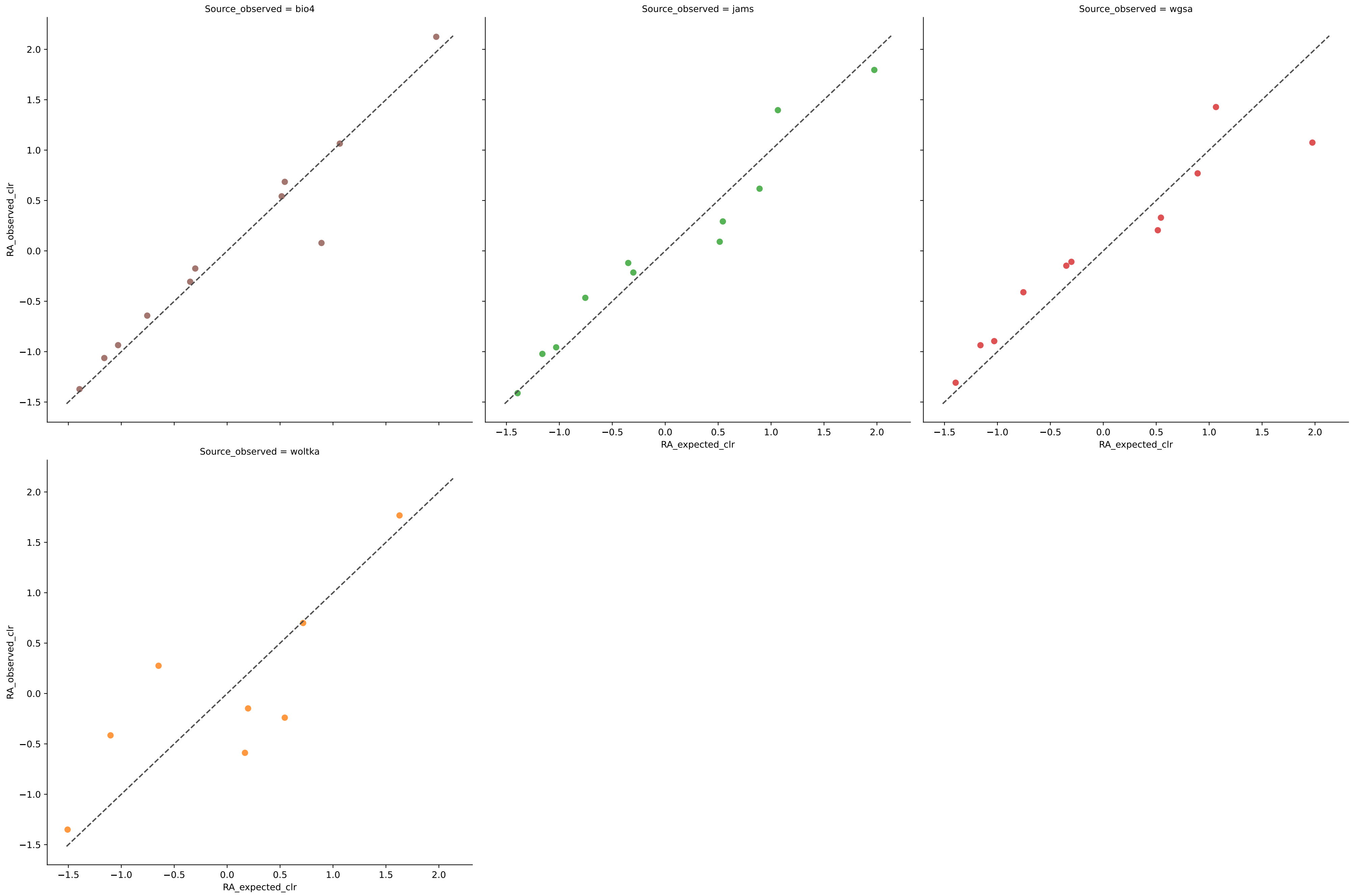


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	13	0.9958	0.0051	0.8159	0.9668	0.0092	100.0000	0.0000
jams	12	0.9396	0.0134	0.7201	0.9198	0.0213	100.0000	0.0000
wgsa	12	0.6014	0.0289	1.1202	0.8267	0.0543	100.0000	0.0000
woltka	9	0.8393	0.0386	2.1938	0.8265	0.0455	100.0000	0.0000

Bivariate Linear Regression for Sample S2 in Experiment camisimGI (Species at filter threshold 0.01)

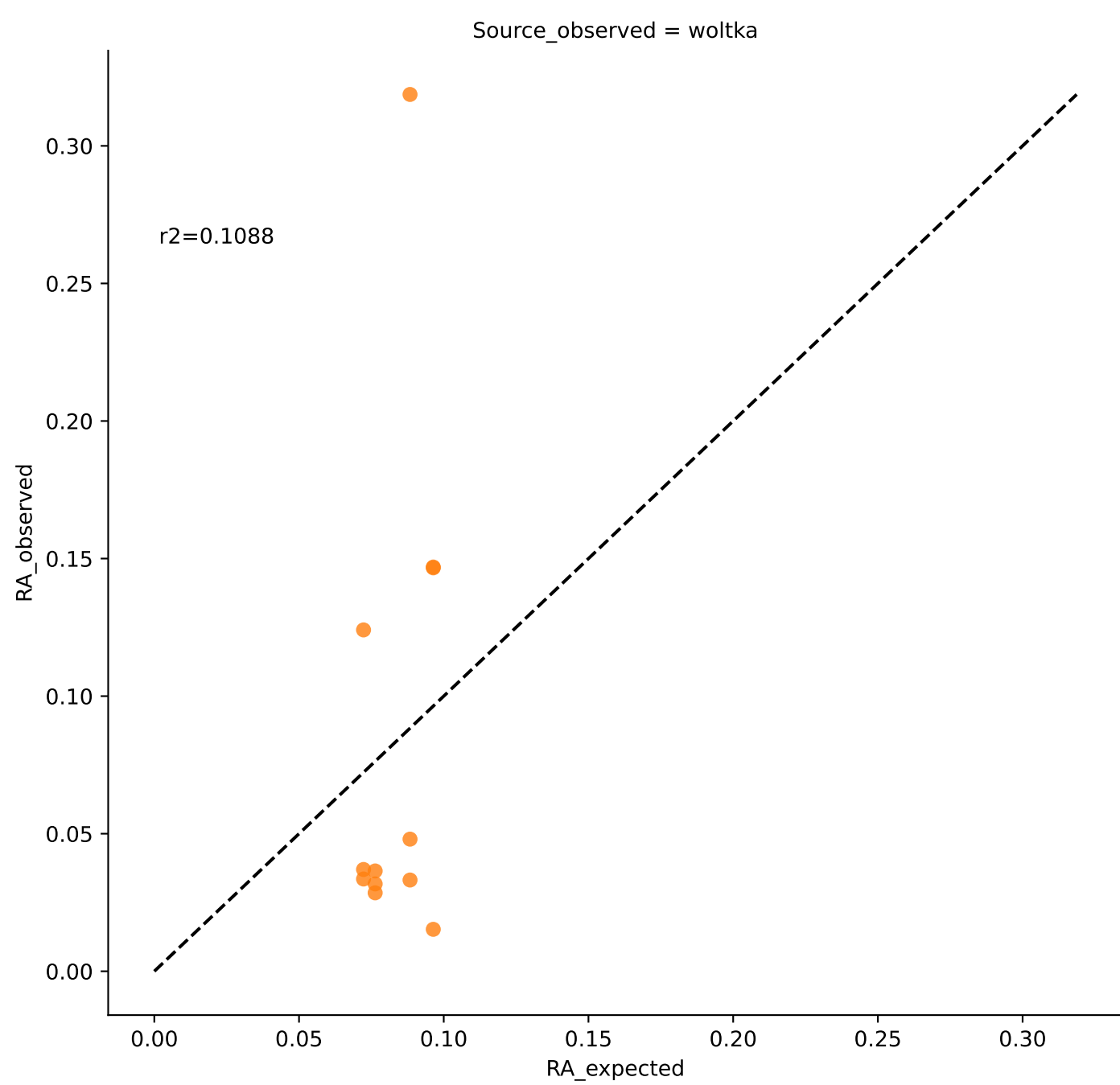
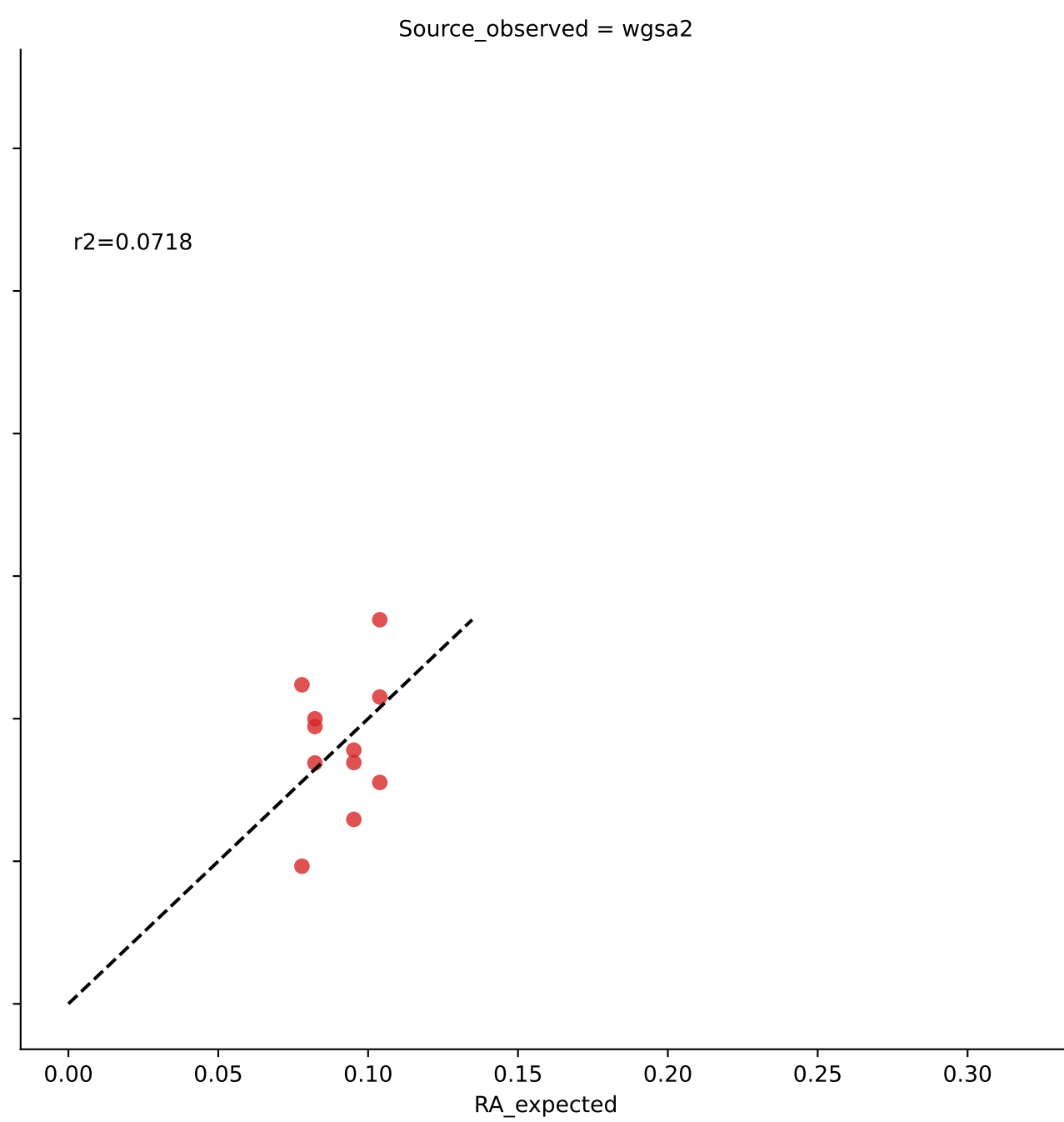
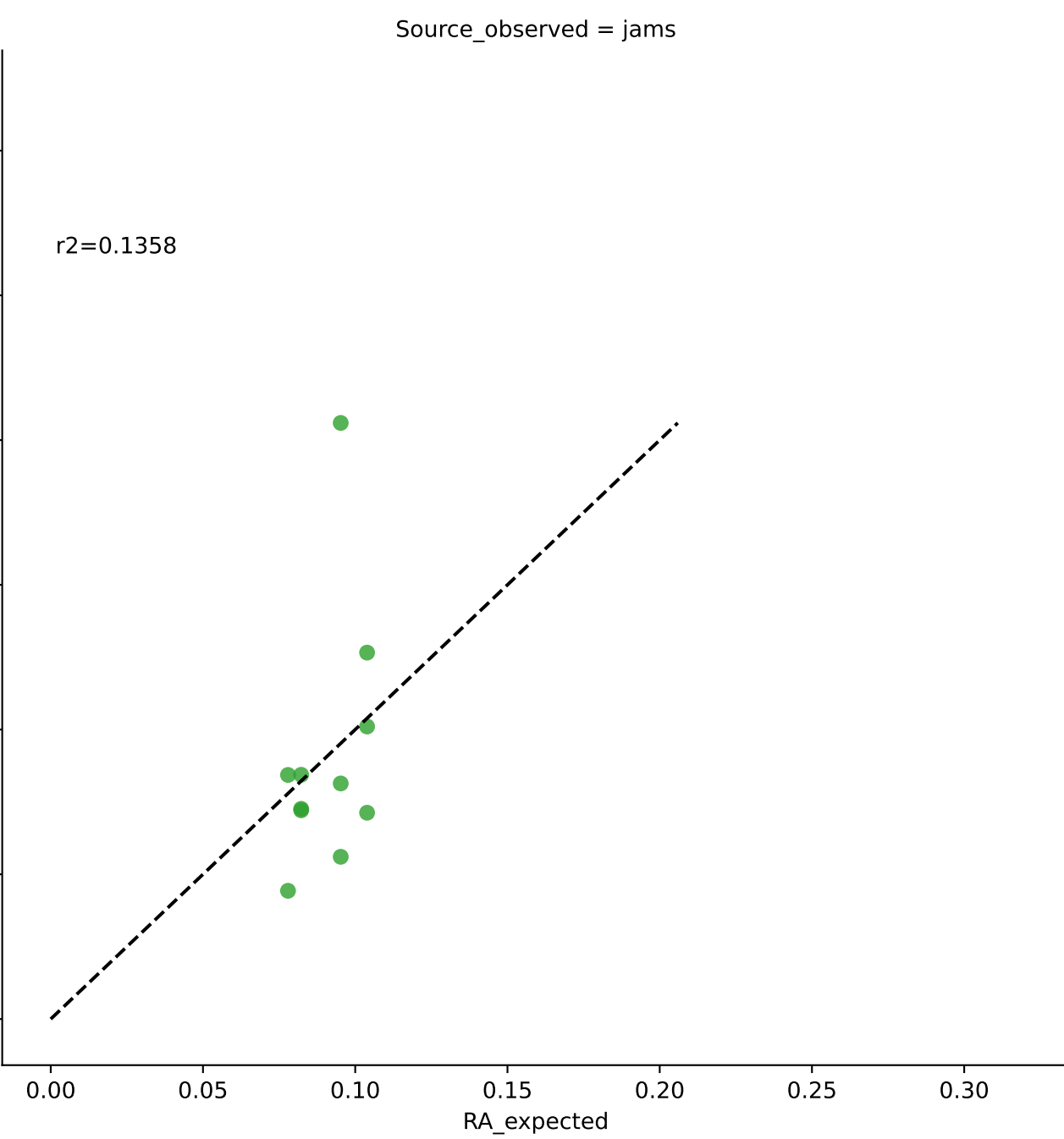
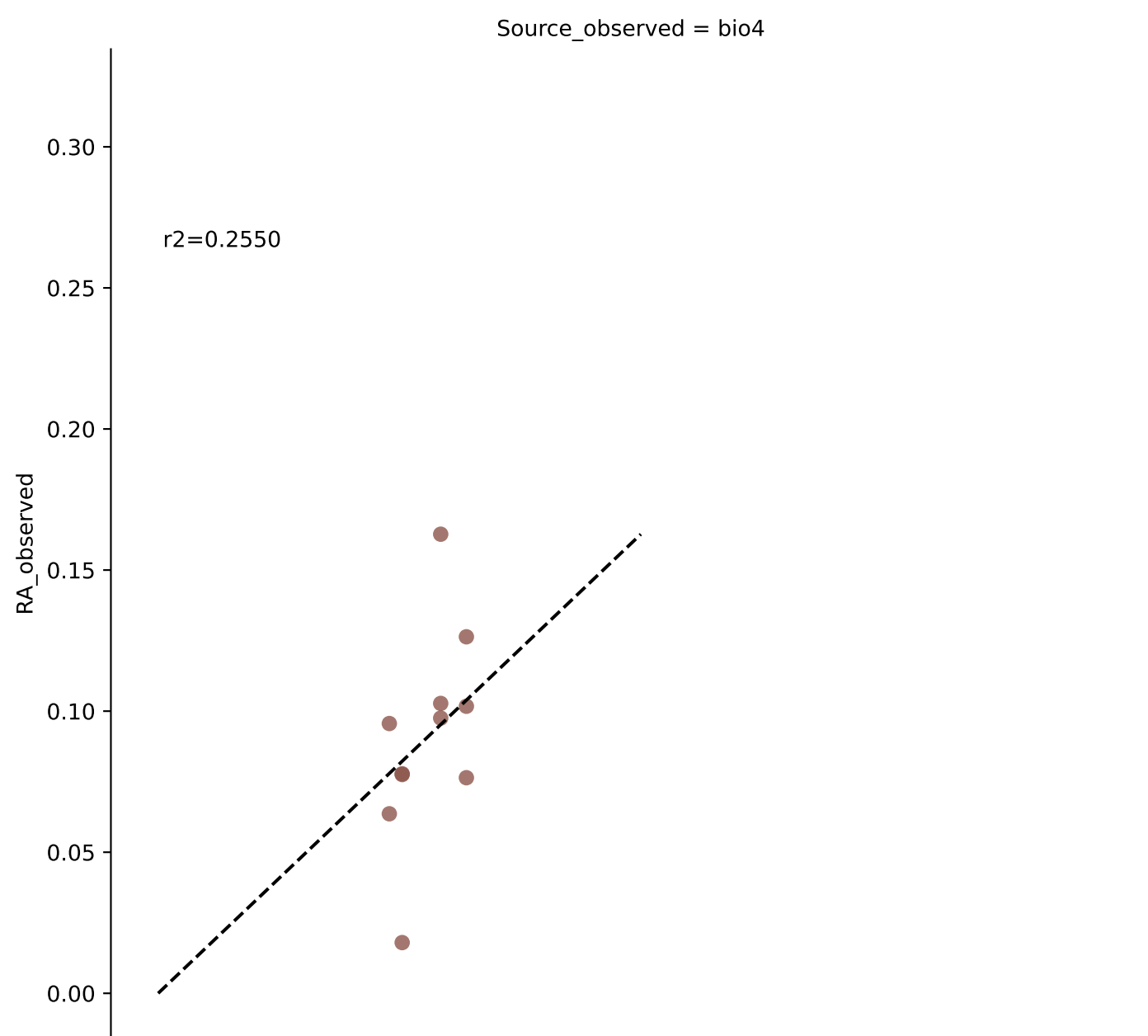


Bivariate Linear Regression for Sample S2 in Experiment camisimGI (Species at filter threshold 0.01)

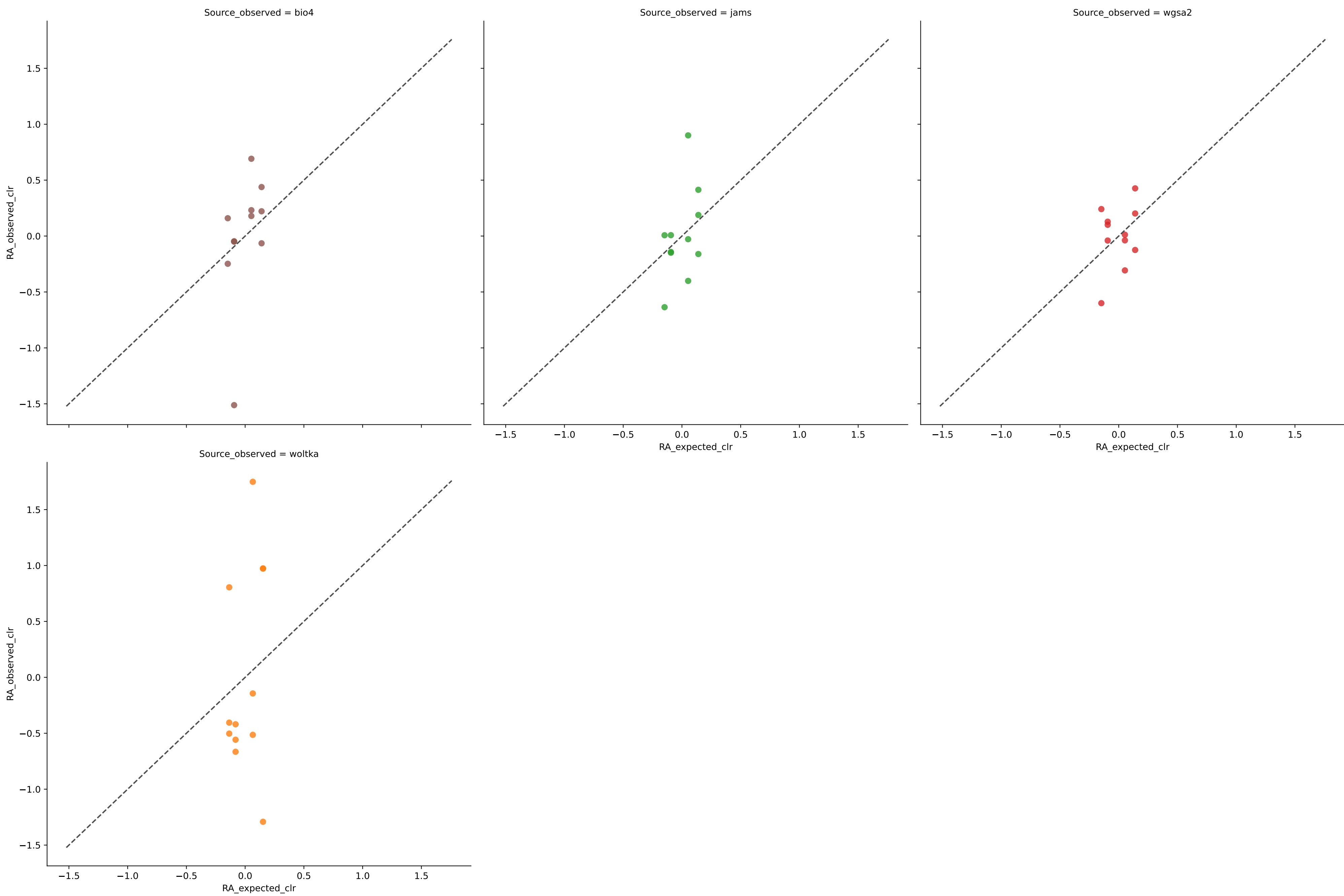


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	11	0.9521	0.0139	0.8675	0.9237	0.0276	100.0000	0.0000
jams	11	0.9212	0.0202	0.7936	0.8891	0.0291	100.0000	0.0000
wgsa	11	0.5558	0.0371	1.1716	0.7961	0.0690	100.0000	0.0000
woltka	8	0.8897	0.0394	1.6356	0.8426	0.0474	100.0000	0.0000

Bivariate Linear Regression for Sample EG in Experiment nist (Genus at filter threshold 0.01)

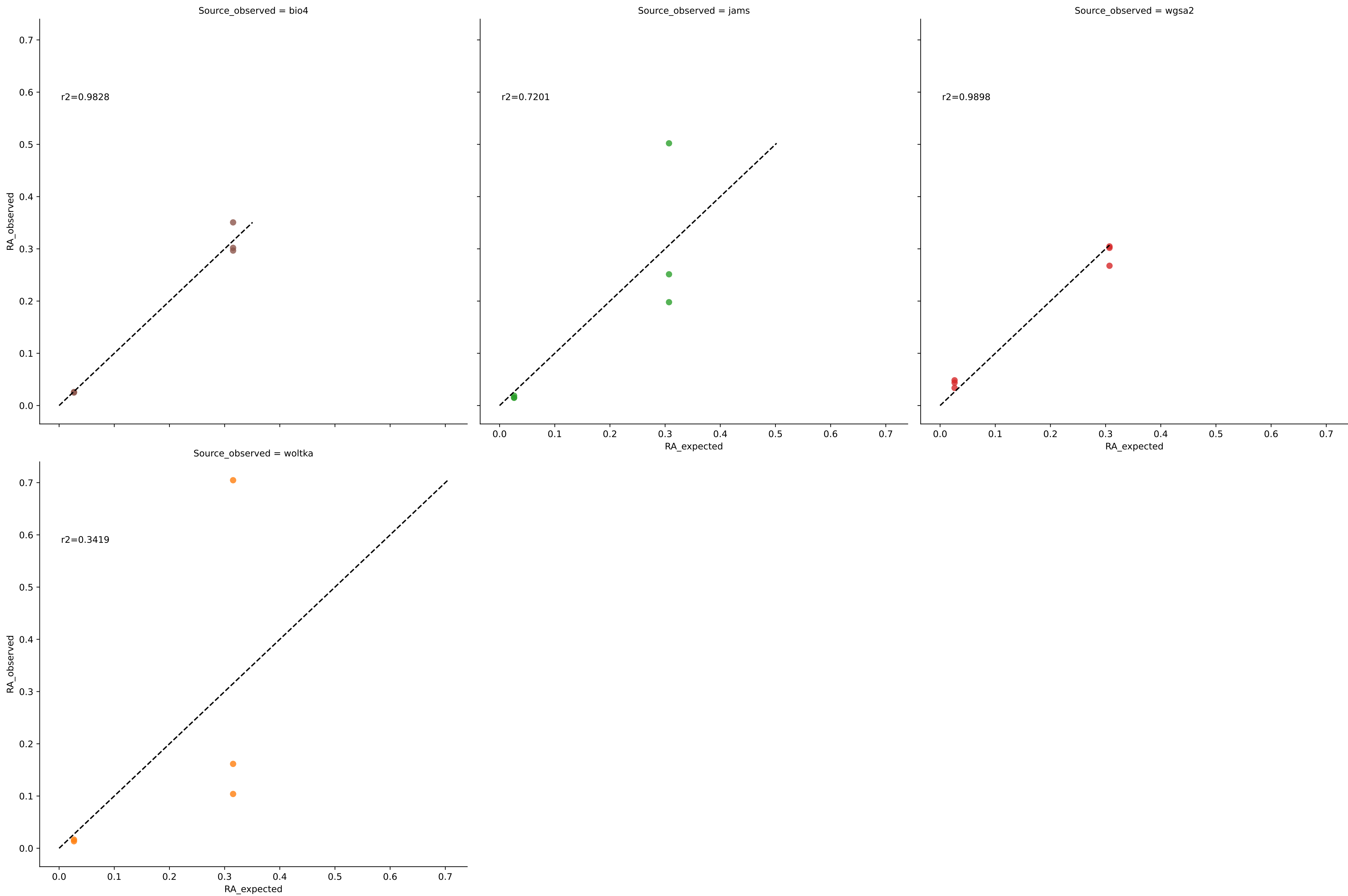


Bivariate Linear Regression for Sample EG in Experiment nist (Genus at filter threshold 0.01)

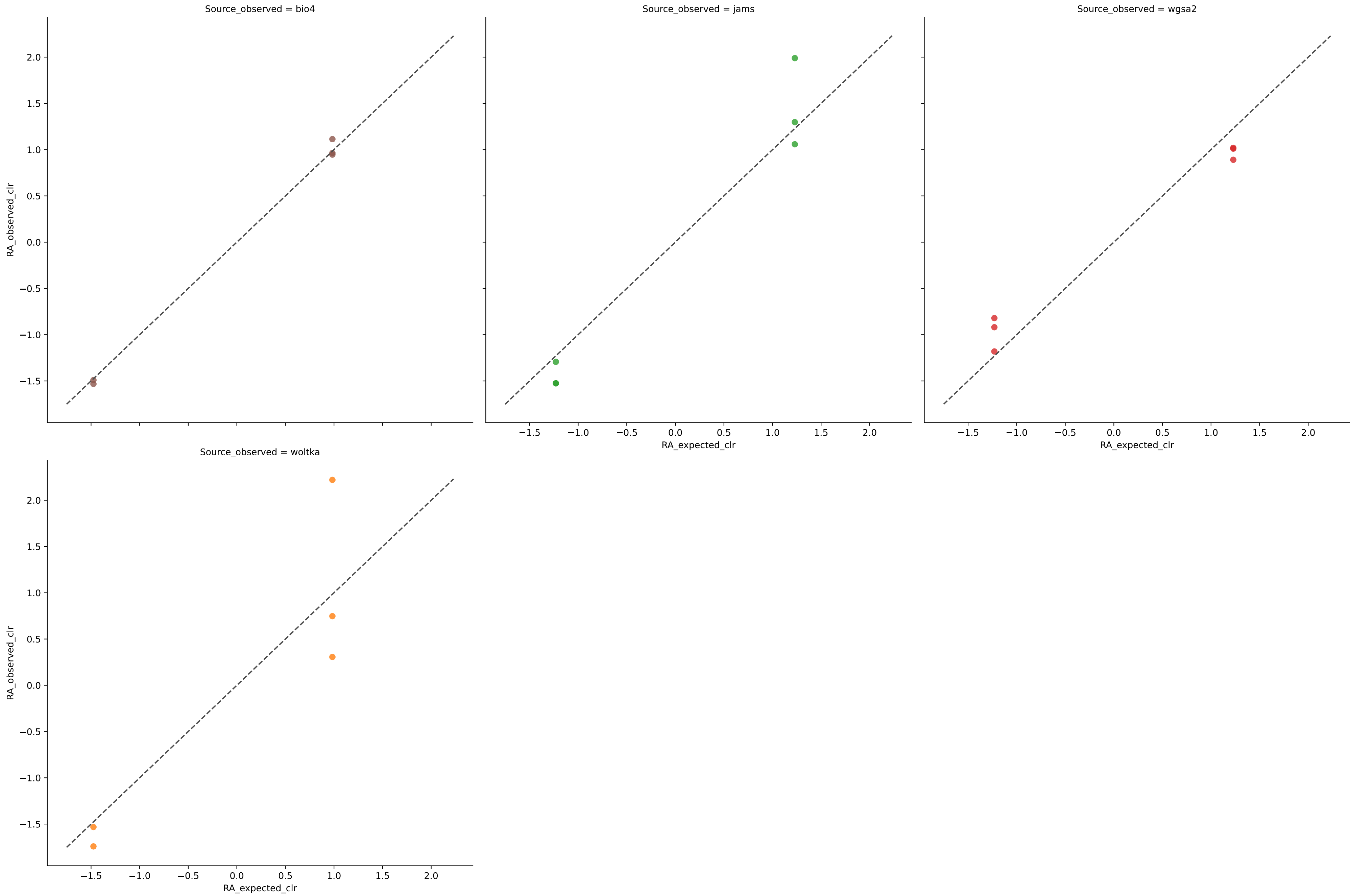


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	11	0.2550	0.0213	1.6469	0.8826	0.0310	100.0000	0.0000
jams	11	0.1358	0.0258	1.1733	0.8581	0.0392	100.0000	0.0000
wgsa2	11	0.0718	0.0188	0.8609	0.8966	0.0220	100.0000	0.0000
woltka	12	0.1088	0.0638	2.9017	0.6171	0.0820	100.0000	0.0000

Bivariate Linear Regression for Sample MIX-A in Experiment nist (Genus at filter threshold 0.01)

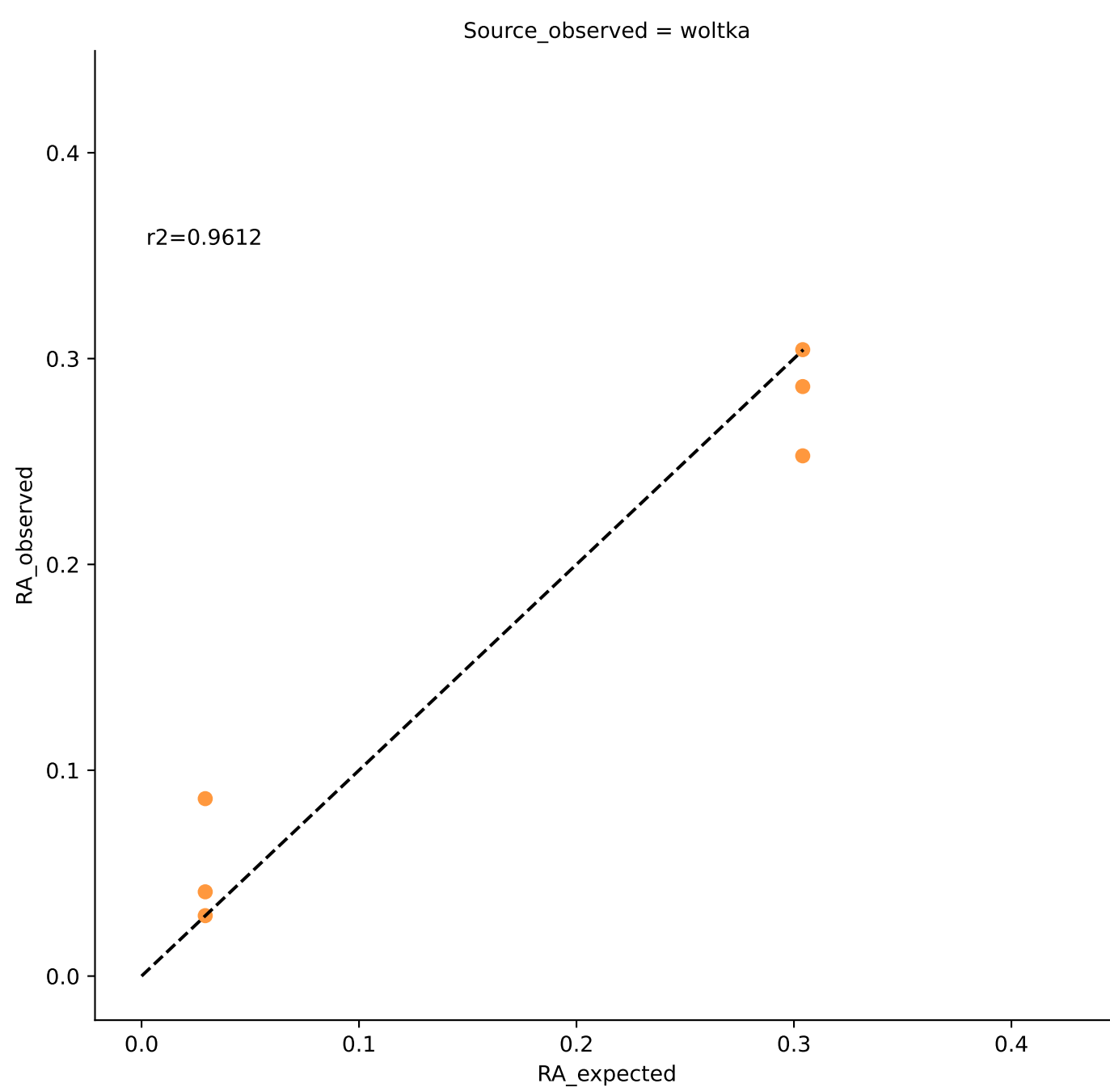
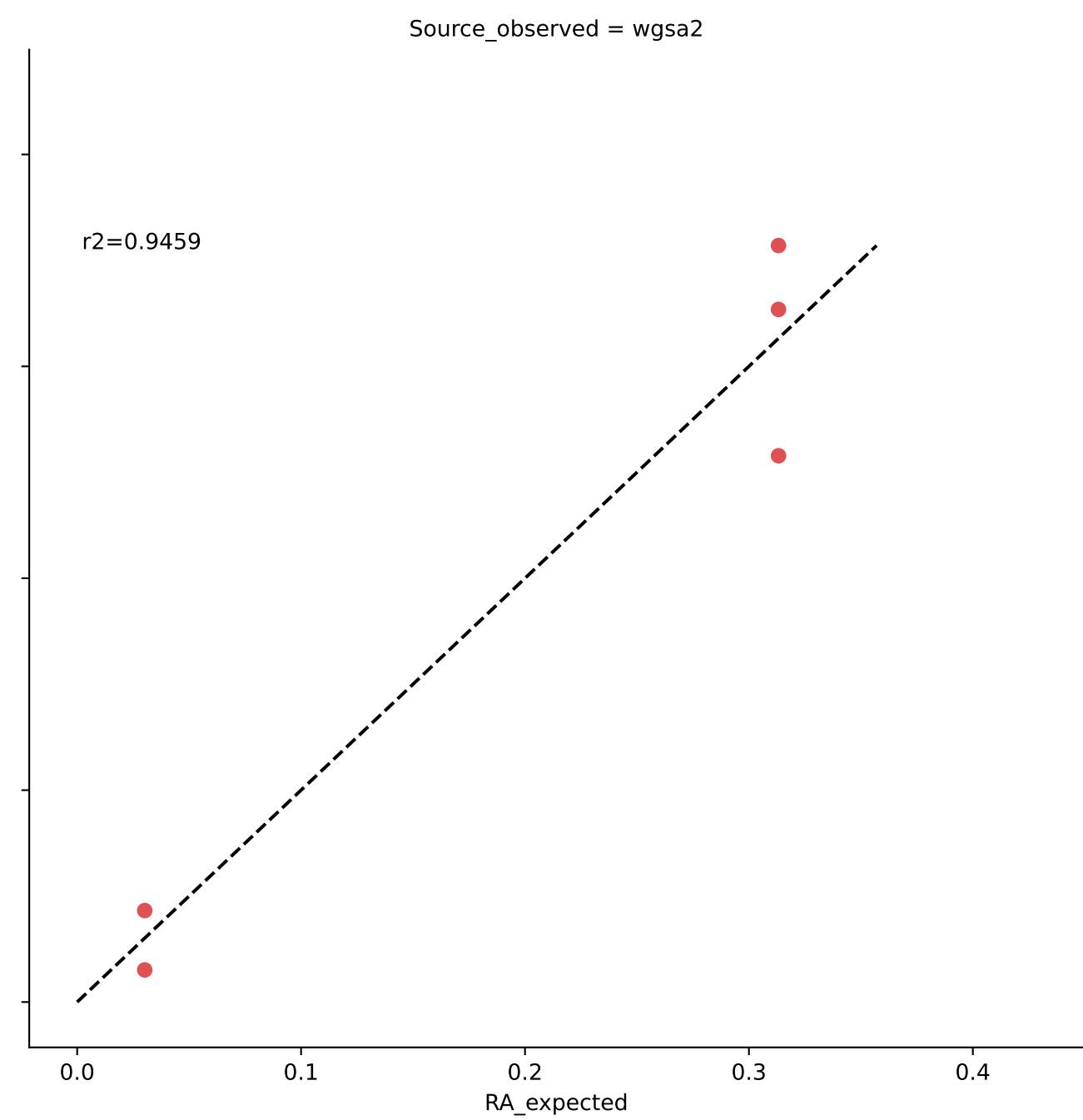
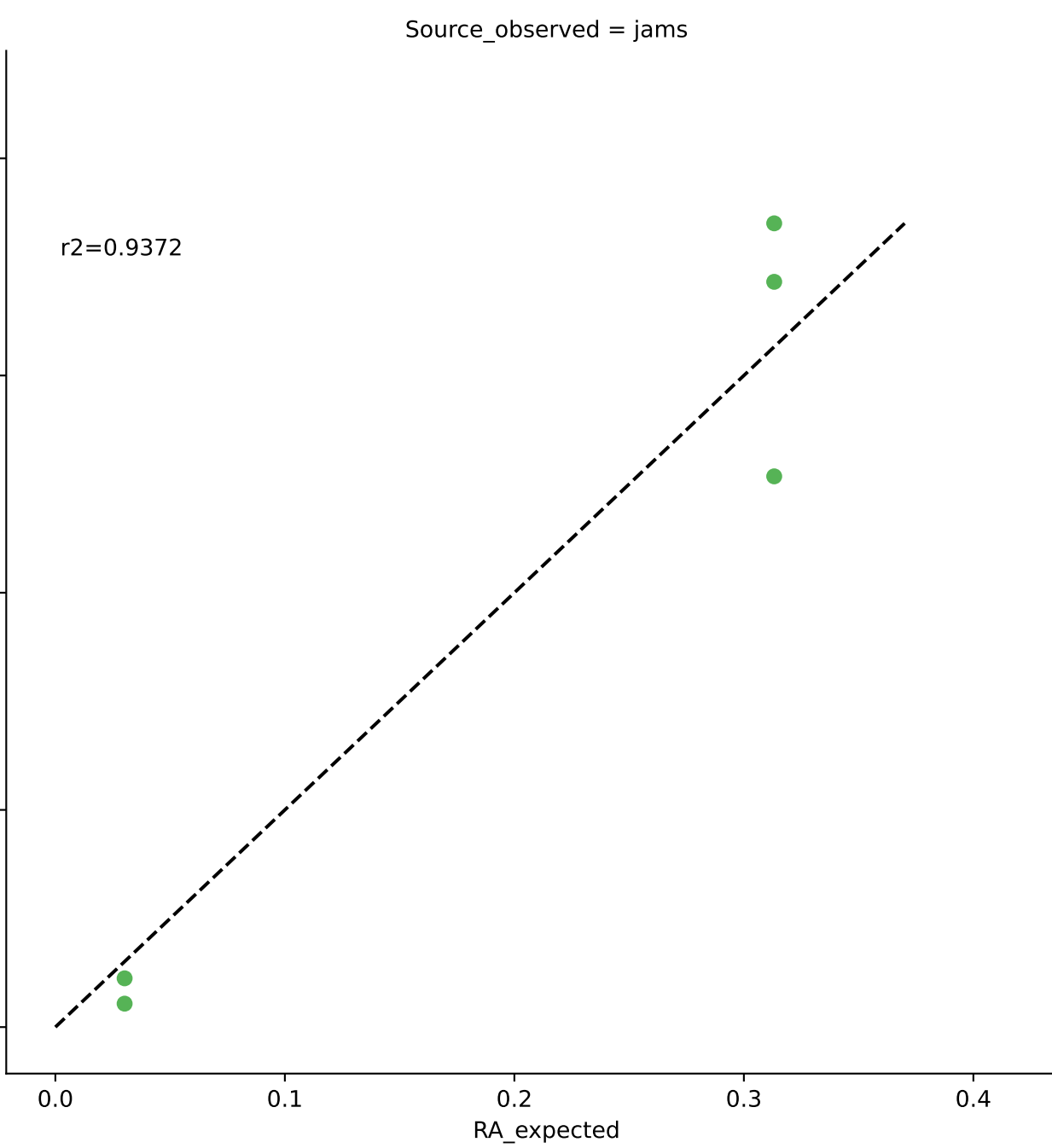
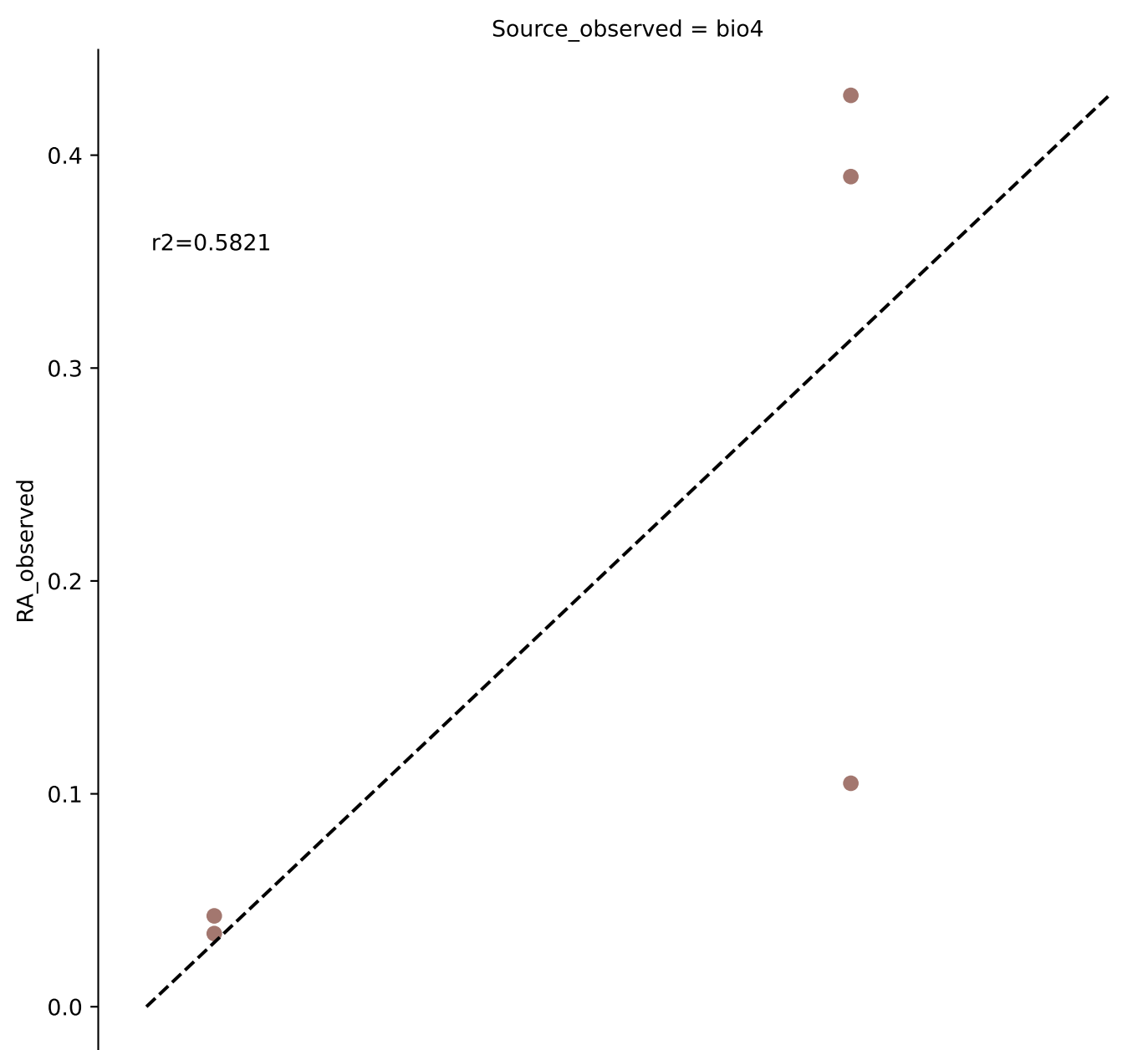


Bivariate Linear Regression for Sample MIX-A in Experiment nist (Genus at filter threshold 0.01)

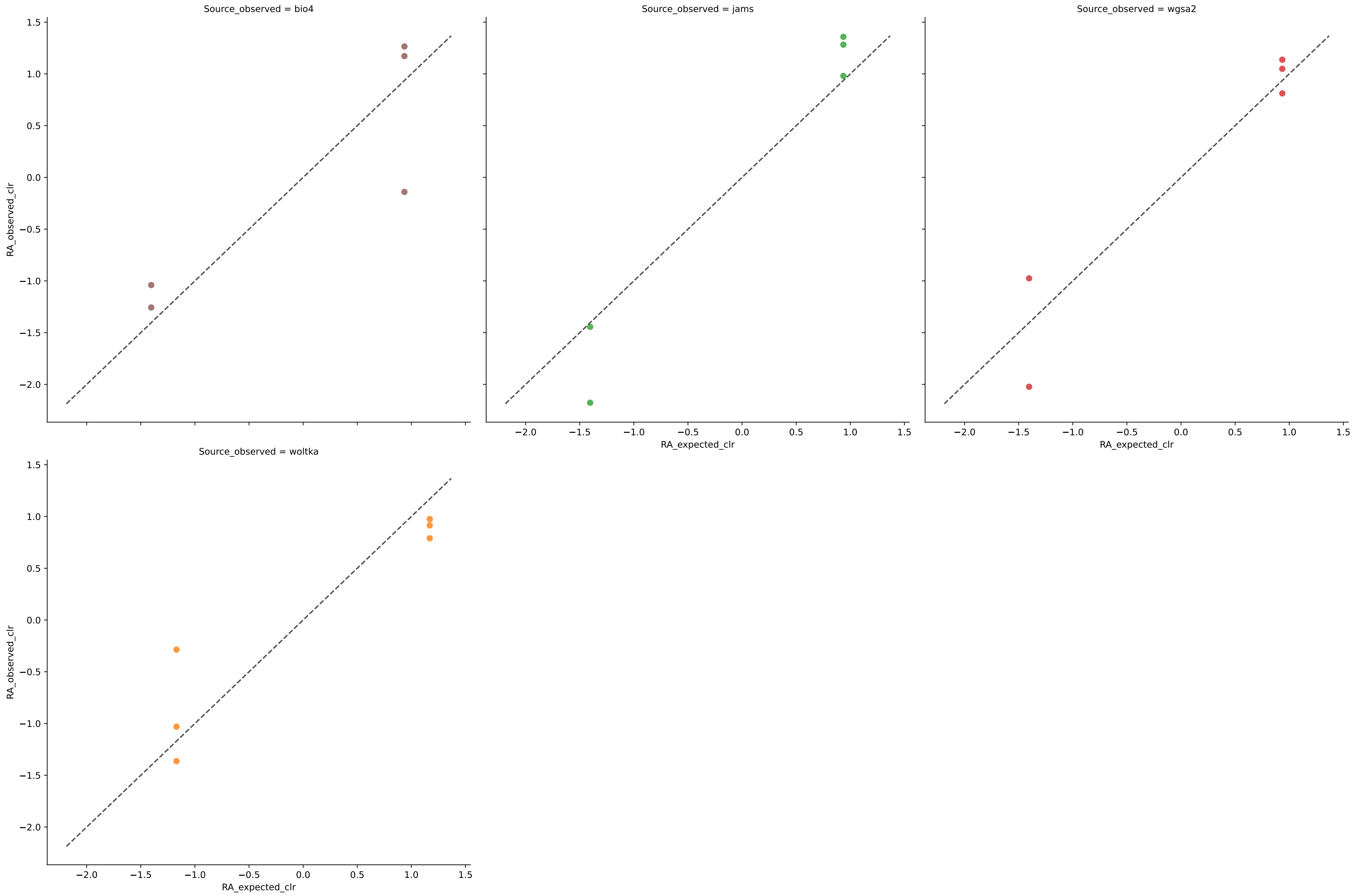


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	5	0.9828	0.0141	0.1490	0.9647	0.0189	100.0000	0.0000
jams	6	0.7201	0.0650	0.8890	0.8050	0.0943	100.0000	0.0000
wgsa2	6	0.9898	0.0157	0.6871	0.9529	0.0202	100.0000	0.0000
woltka	5	0.3419	0.1557	1.4554	0.6107	0.2098	100.0000	0.0000

Bivariate Linear Regression for Sample MIX-B in Experiment nist (Genus at filter threshold 0.01)

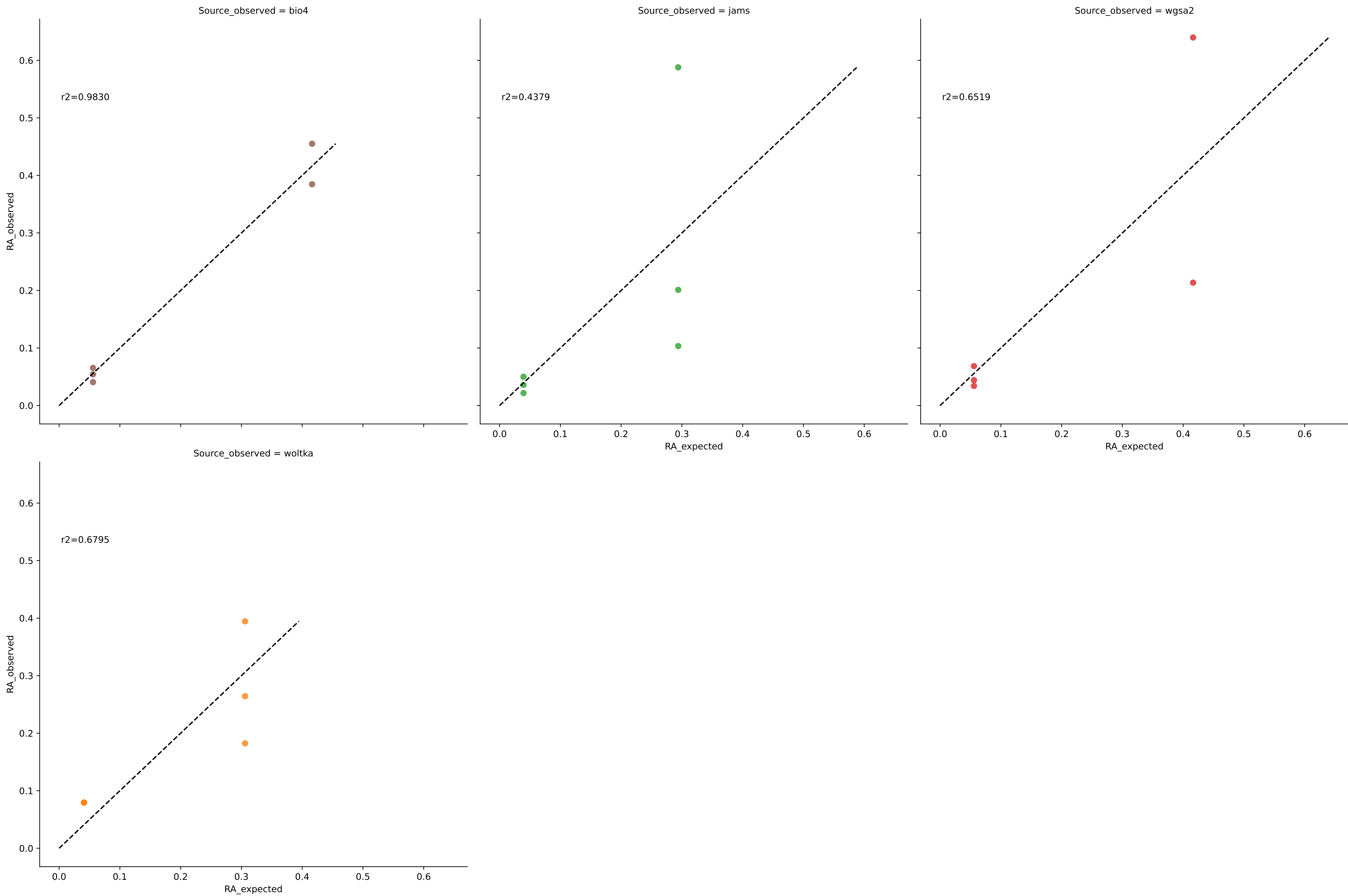


Bivariate Linear Regression for Sample MIX-B in Experiment nist (Genus at filter threshold 0.01)

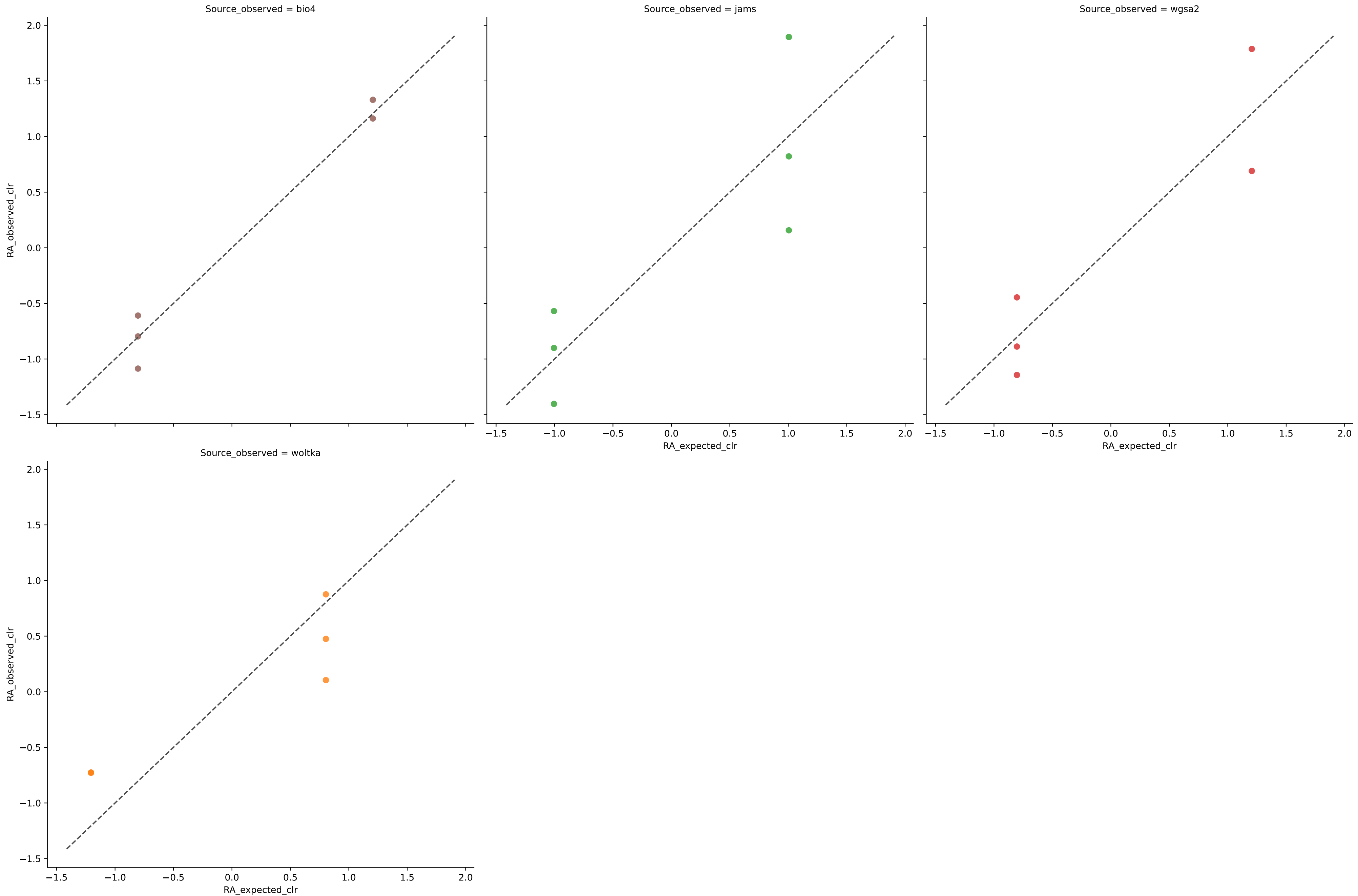


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	5	0.5821	0.0833	1.2151	0.7917	0.1119	100.0000	0.0000
jams	5	0.9372	0.0347	0.9485	0.9132	0.0403	100.0000	0.0000
wgsa2	5	0.9459	0.0282	0.7972	0.9296	0.0334	100.0000	0.0000
woltka	6	0.9612	0.0230	1.0419	0.9311	0.0324	100.0000	0.0000

Bivariate Linear Regression for Sample MIX-C in Experiment nist (Genus at filter threshold 0.01)

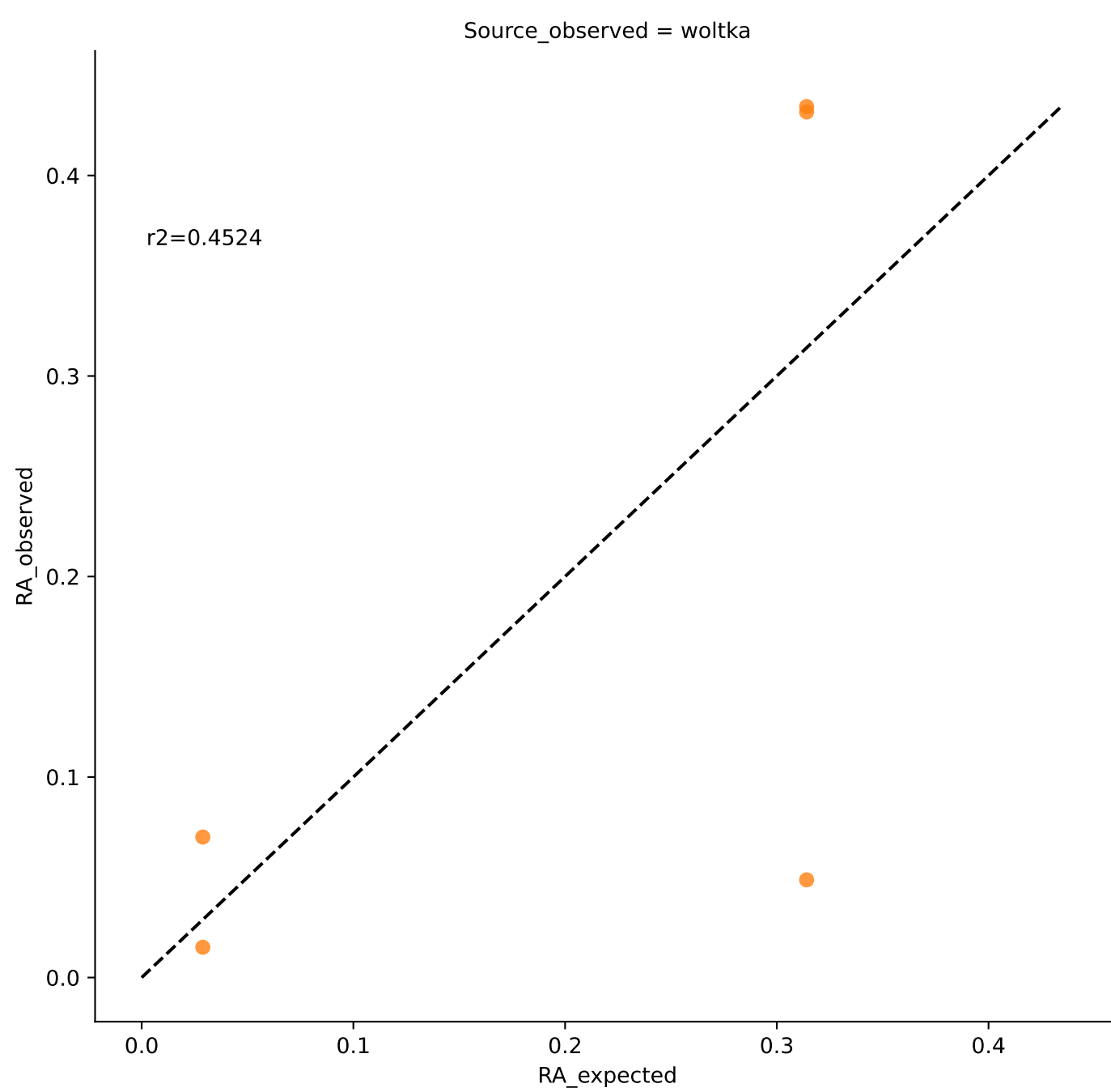
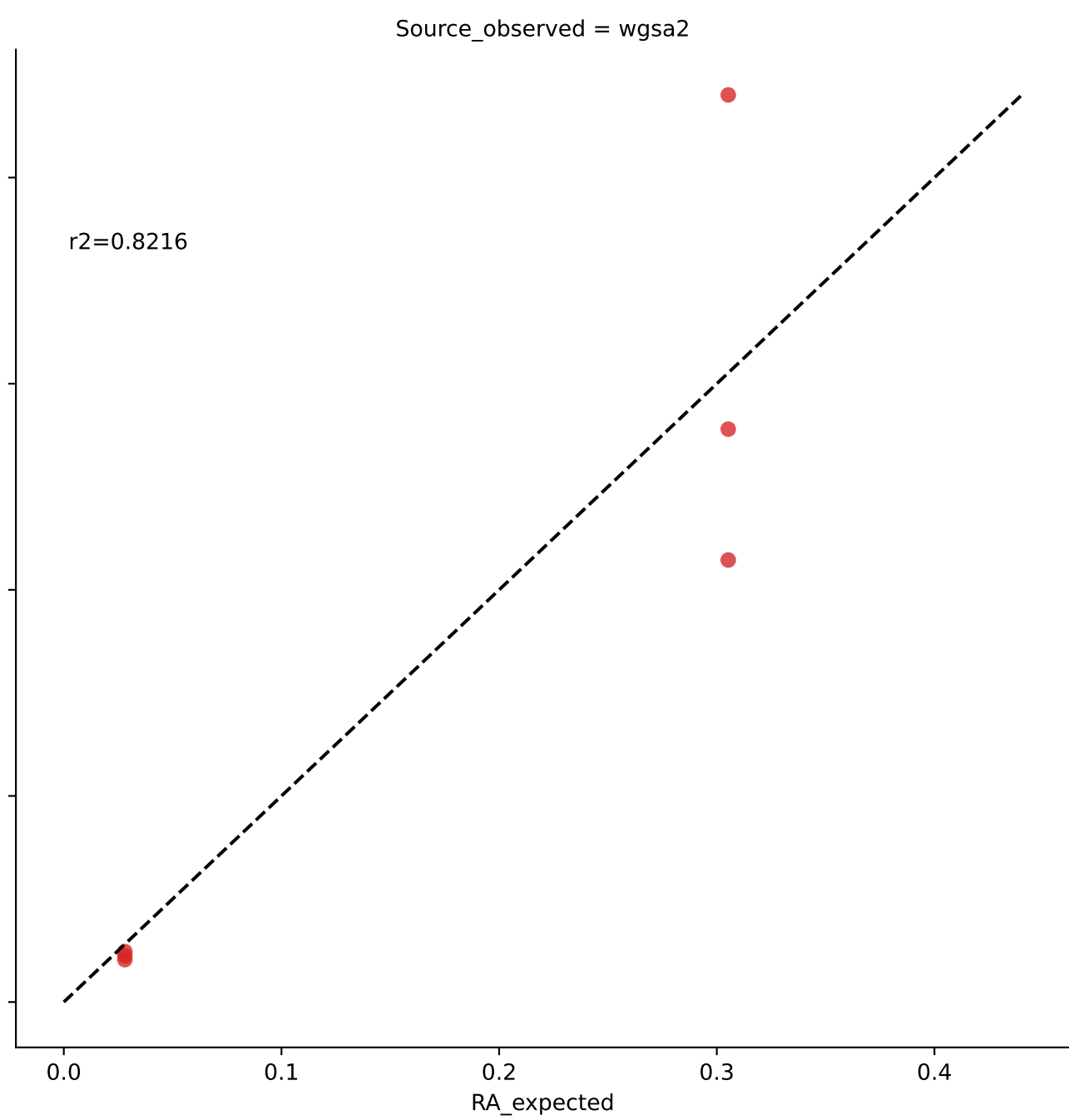
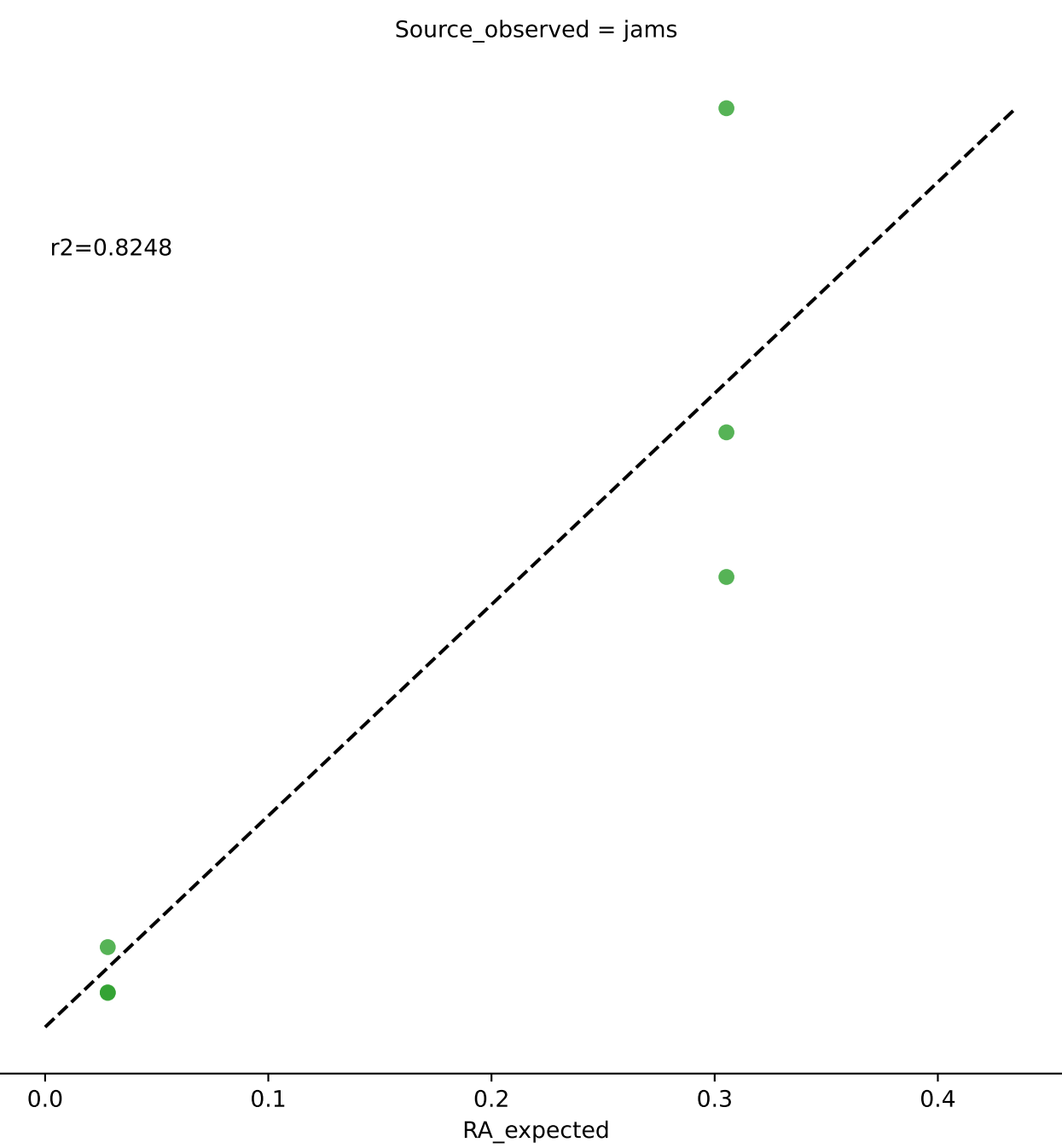
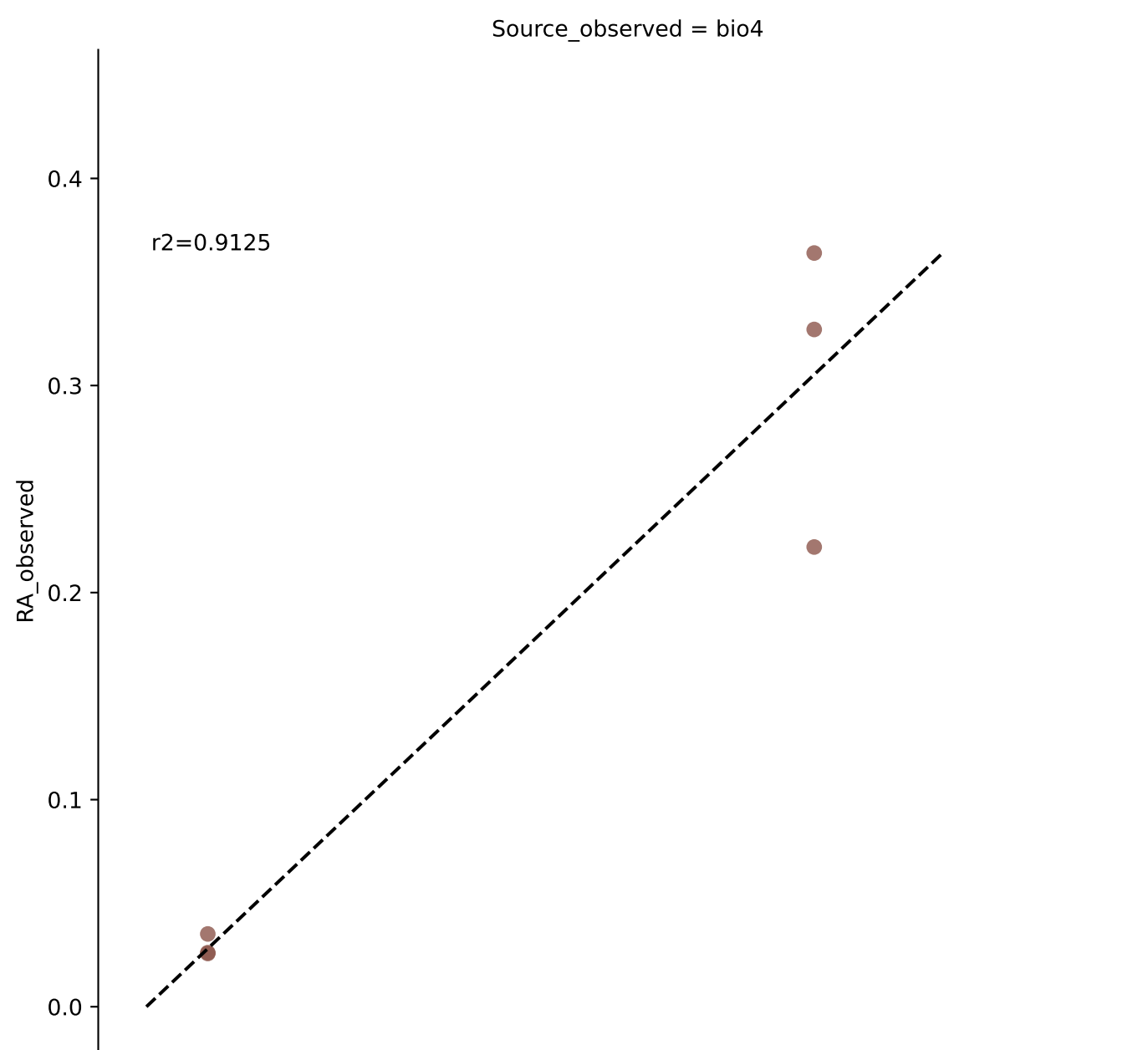


Bivariate Linear Regression for Sample MIX-C in Experiment nist (Genus at filter threshold 0.01)

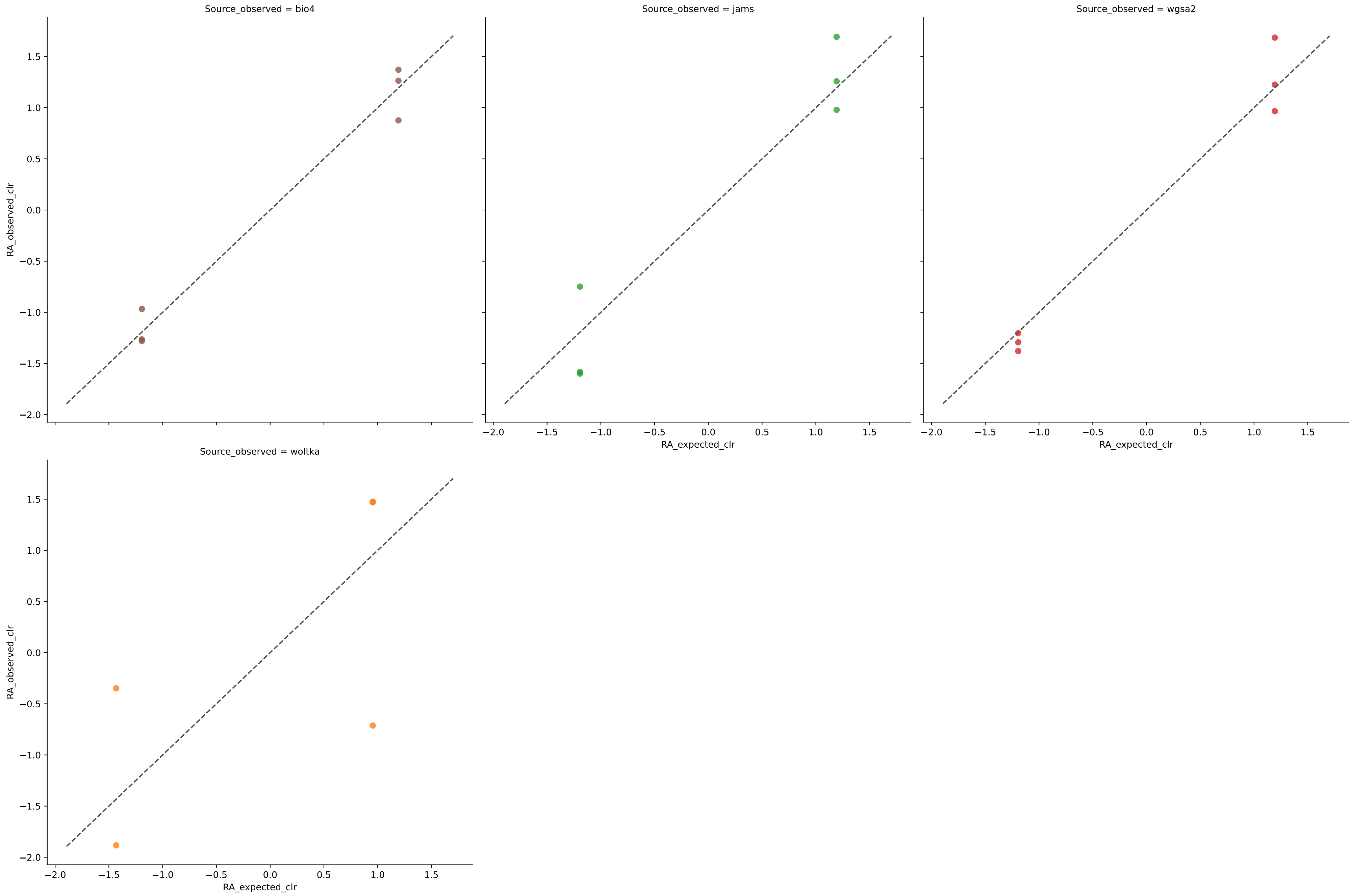


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	5	0.9830	0.0194	0.3674	0.9516	0.0238	100.0000	0.0000
jams	6	0.4379	0.1015	1.3797	0.6954	0.1482	100.0000	0.0000
wgsa2	5	0.6519	0.0945	0.9244	0.7639	0.1355	100.0000	0.0000
woltka	5	0.6795	0.0661	1.0297	0.8348	0.0745	100.0000	0.0000

Bivariate Linear Regression for Sample MIX-D in Experiment nist (Genus at filter threshold 0.01)

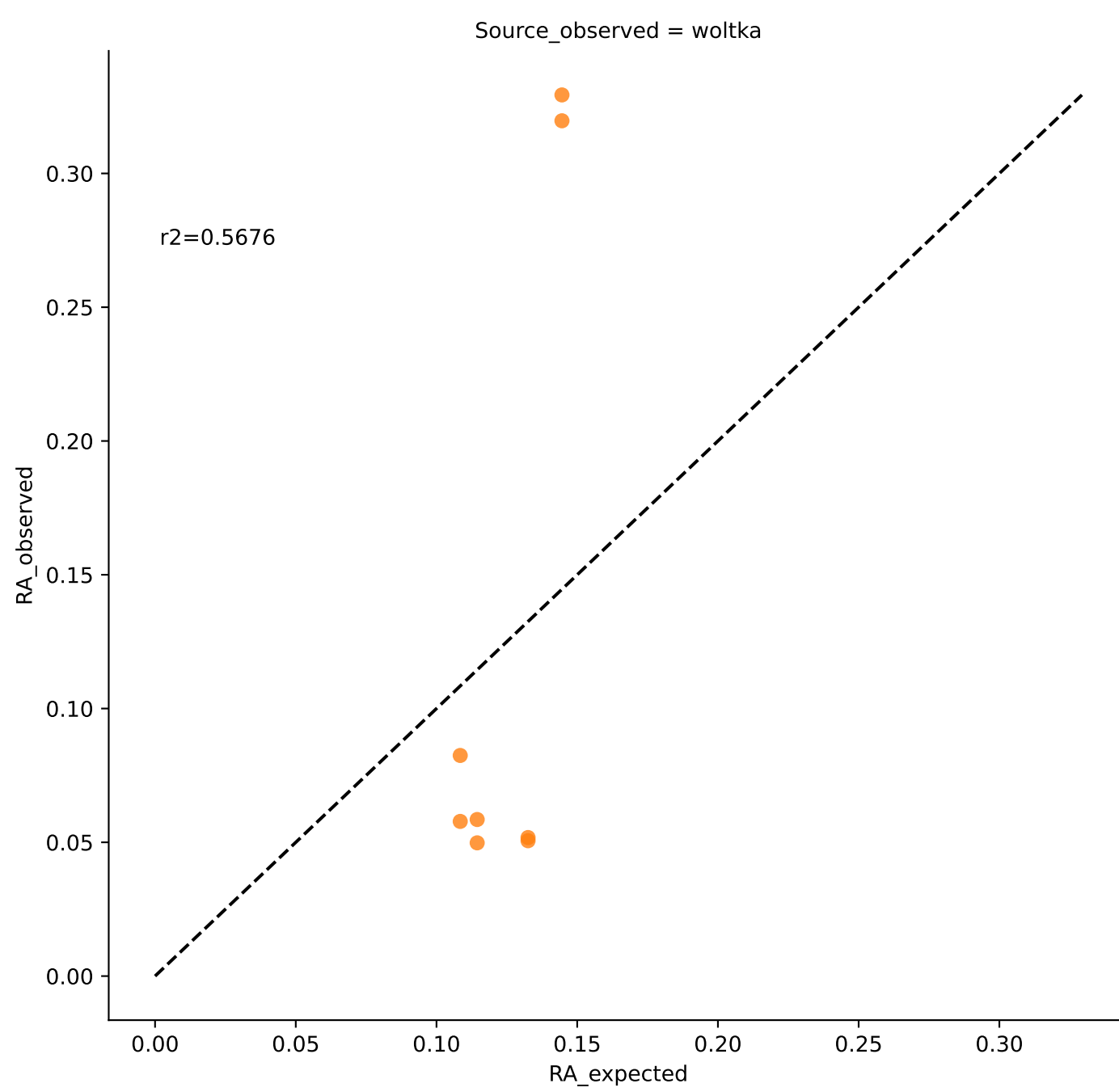
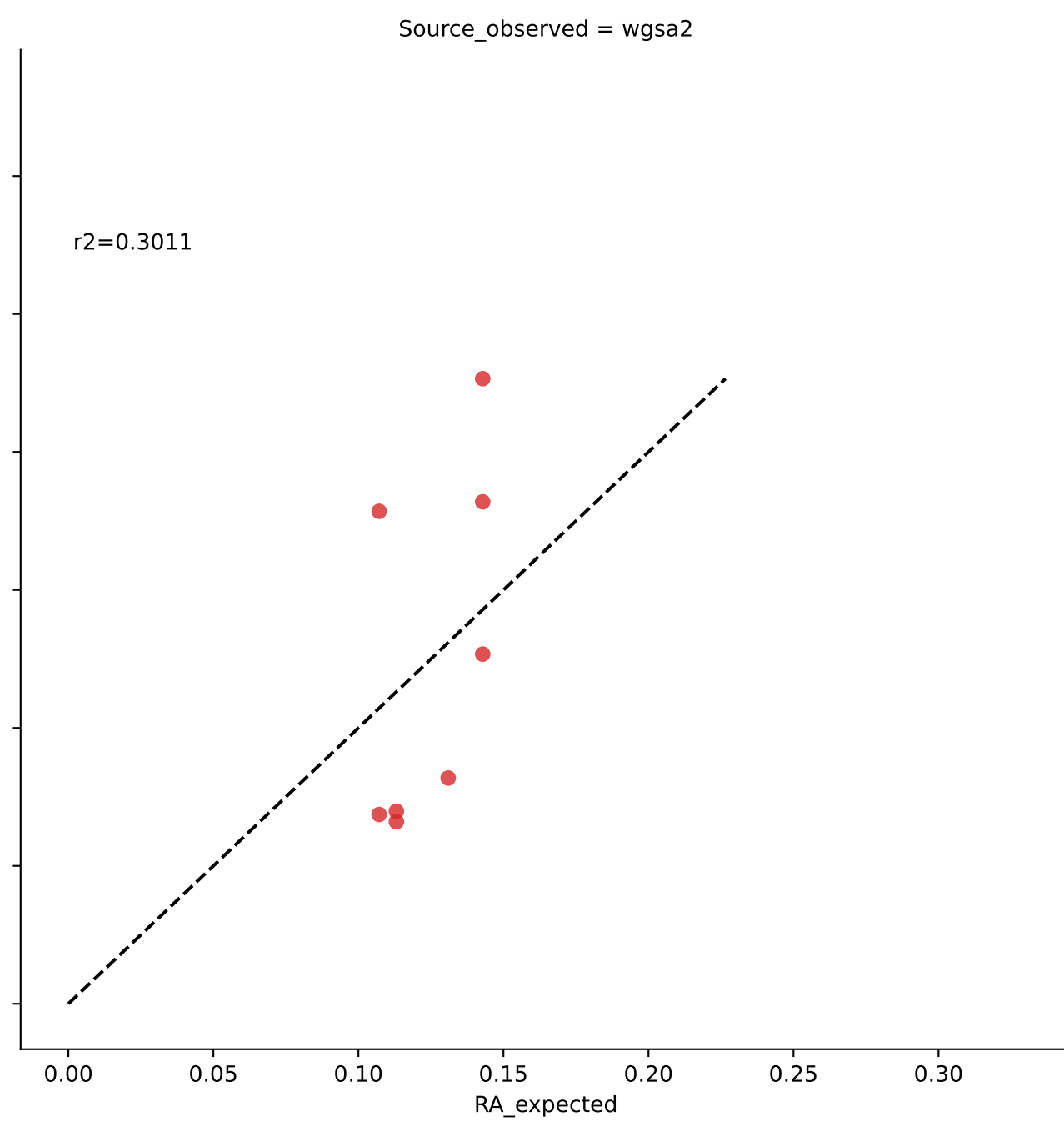
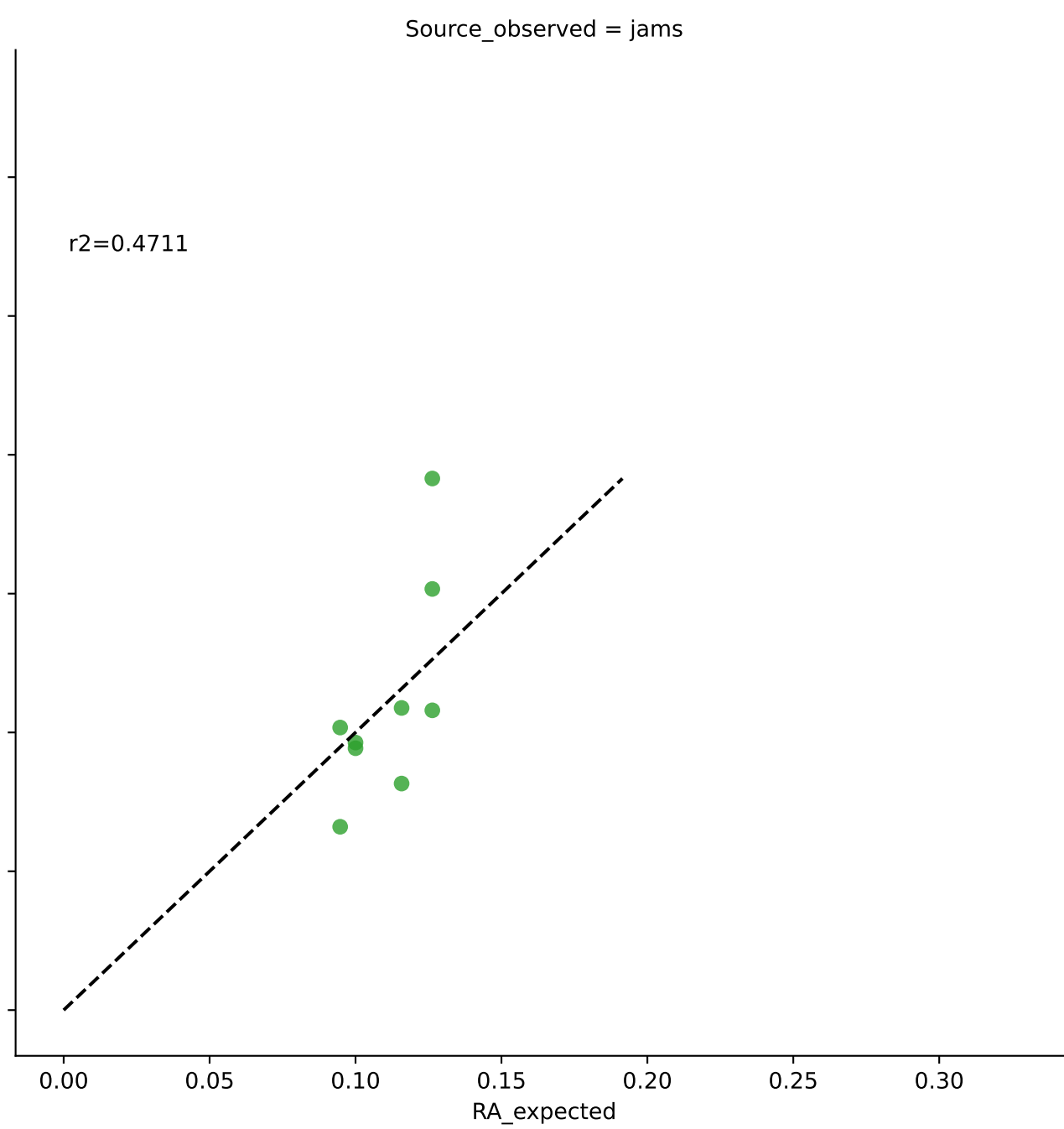
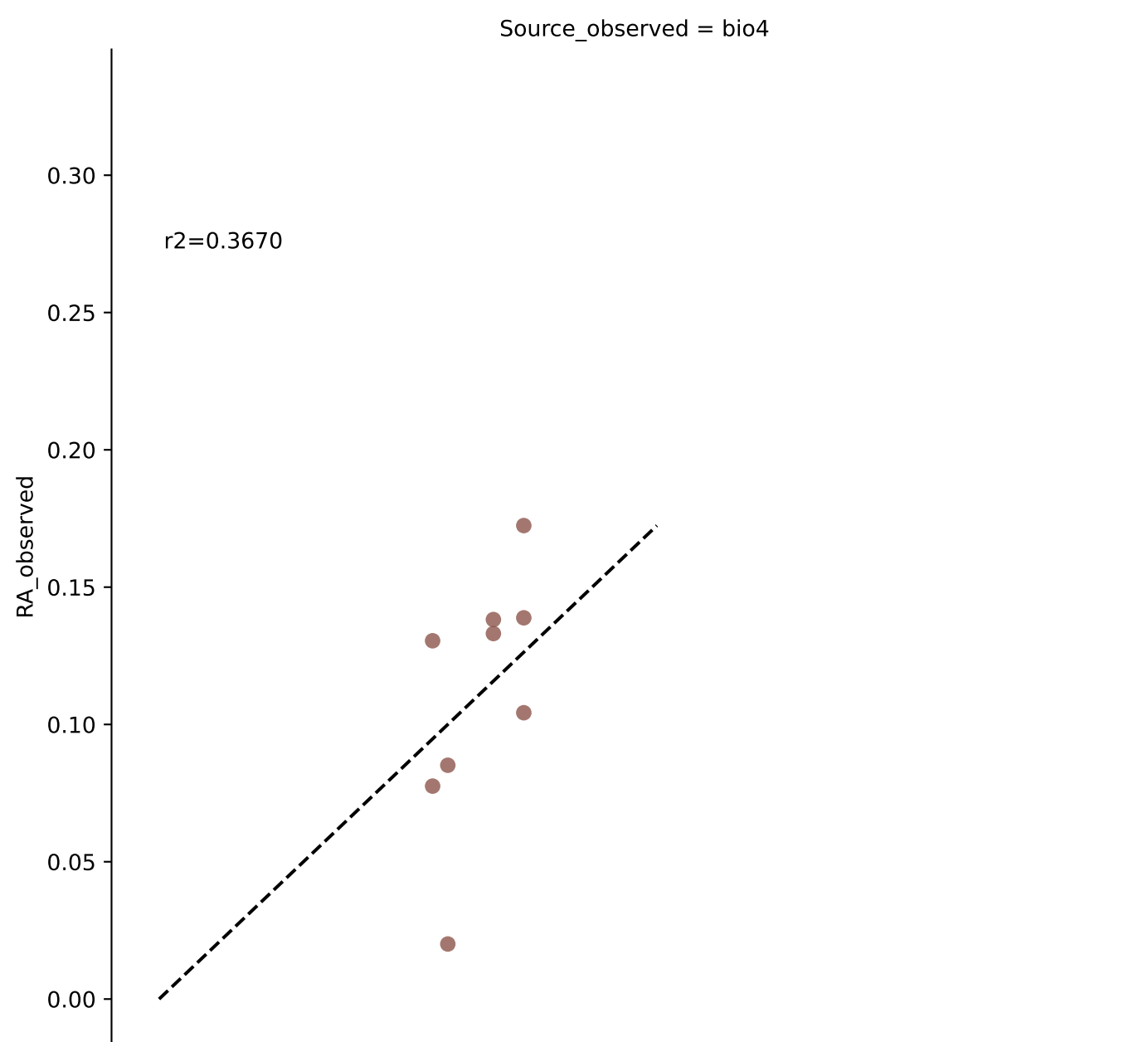


Bivariate Linear Regression for Sample MIX-D in Experiment nist (Genus at filter threshold 0.01)

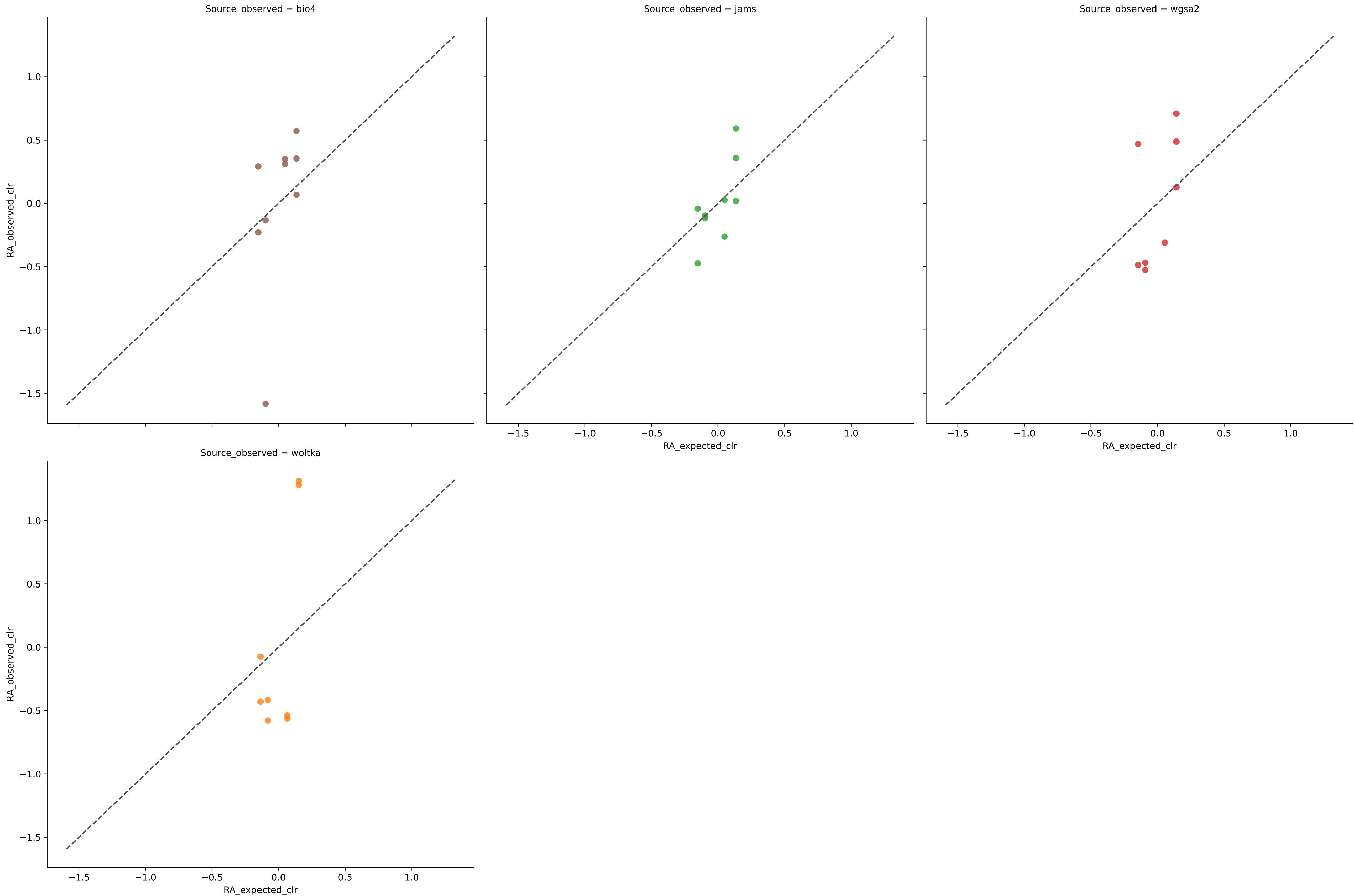


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	6	0.9125	0.0292	0.4479	0.9124	0.0426	100.0000	0.0000
jams	6	0.8248	0.0465	0.9017	0.8605	0.0661	100.0000	0.0000
wgsa2	6	0.8216	0.0449	0.5821	0.8652	0.0674	100.0000	0.0000
woltka	5	0.4524	0.1117	2.1662	0.7209	0.1419	100.0000	0.0000

Bivariate Linear Regression for Sample EG in Experiment nist (Species at filter threshold 0.01)

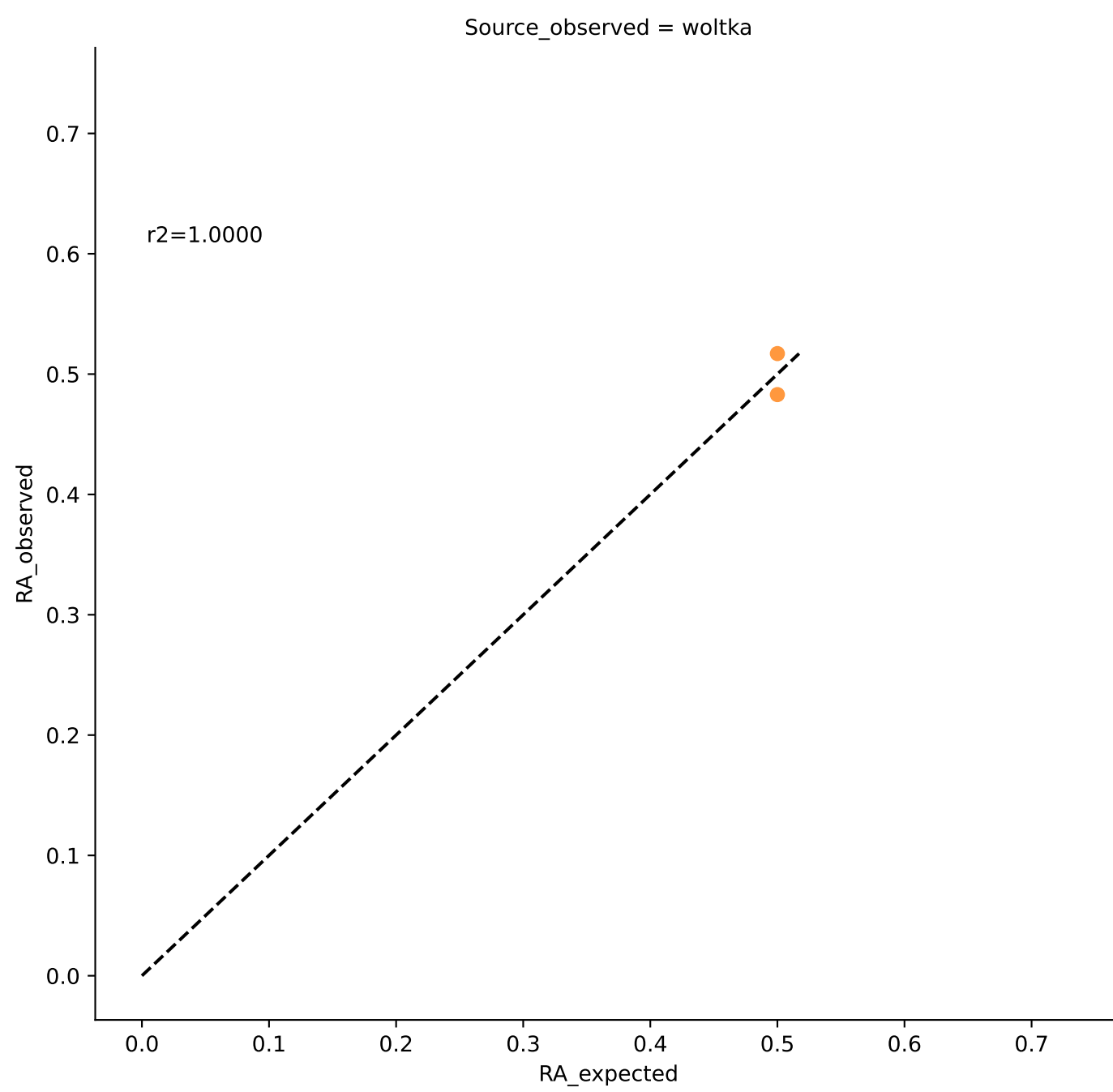
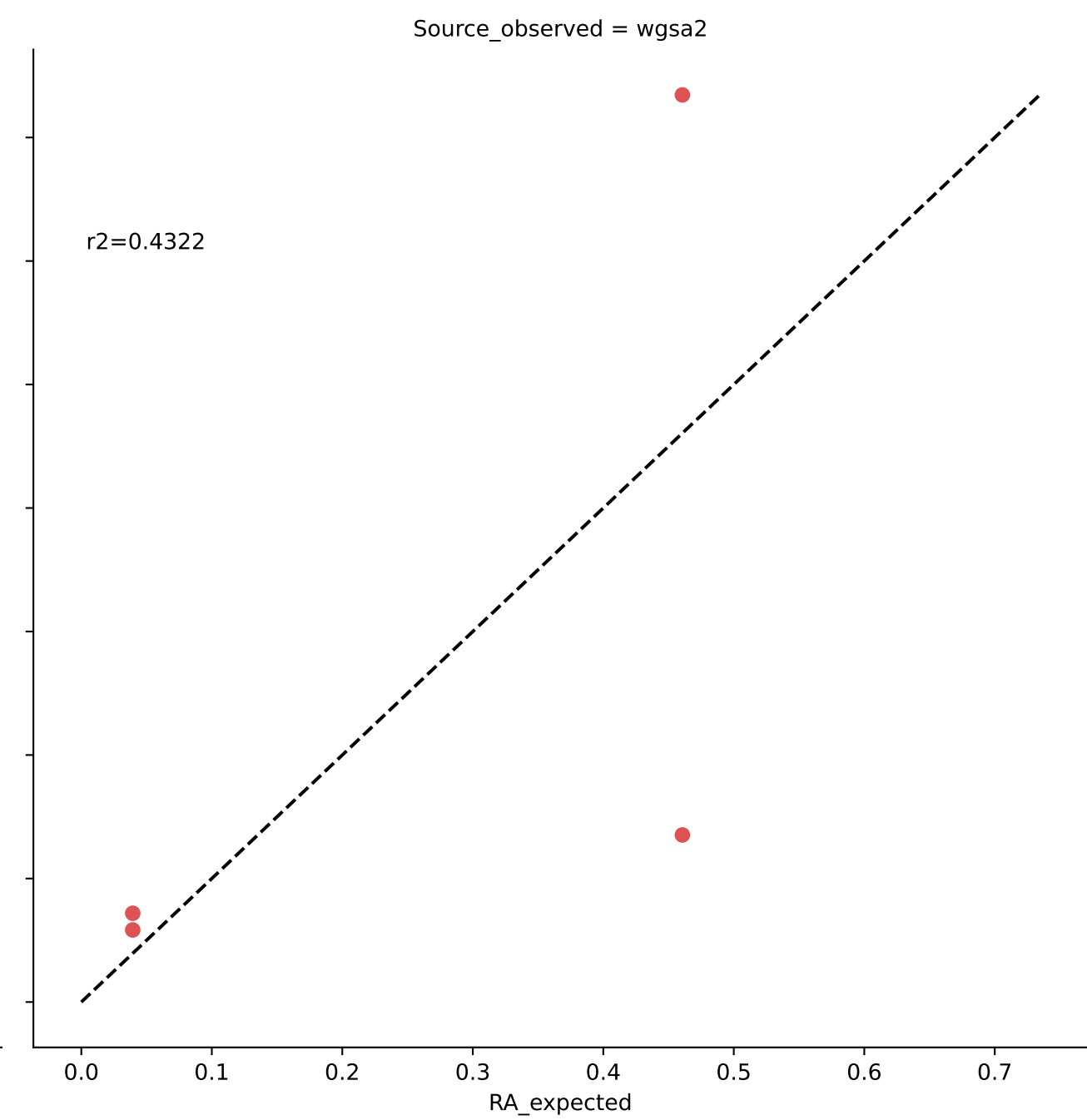
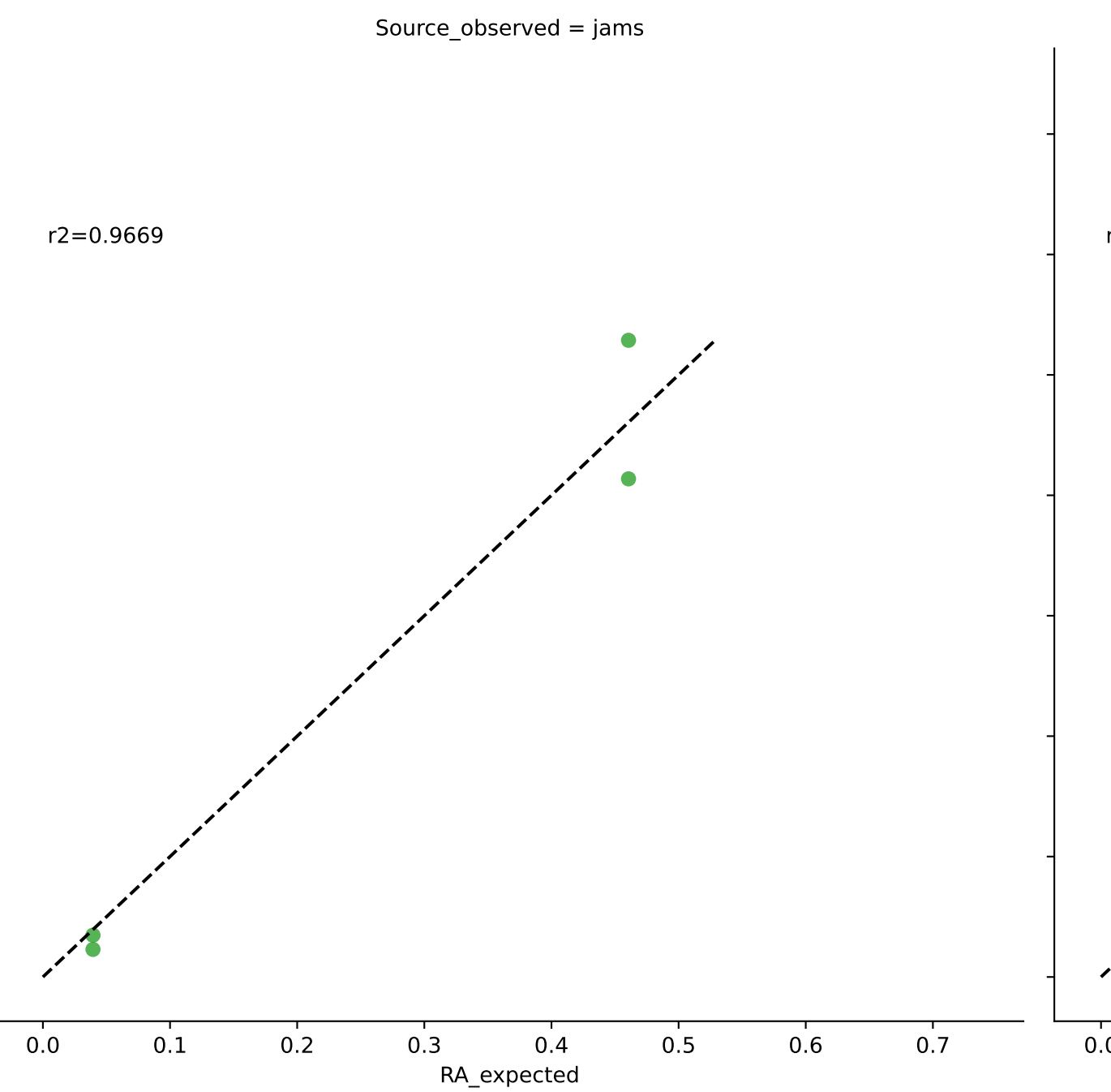
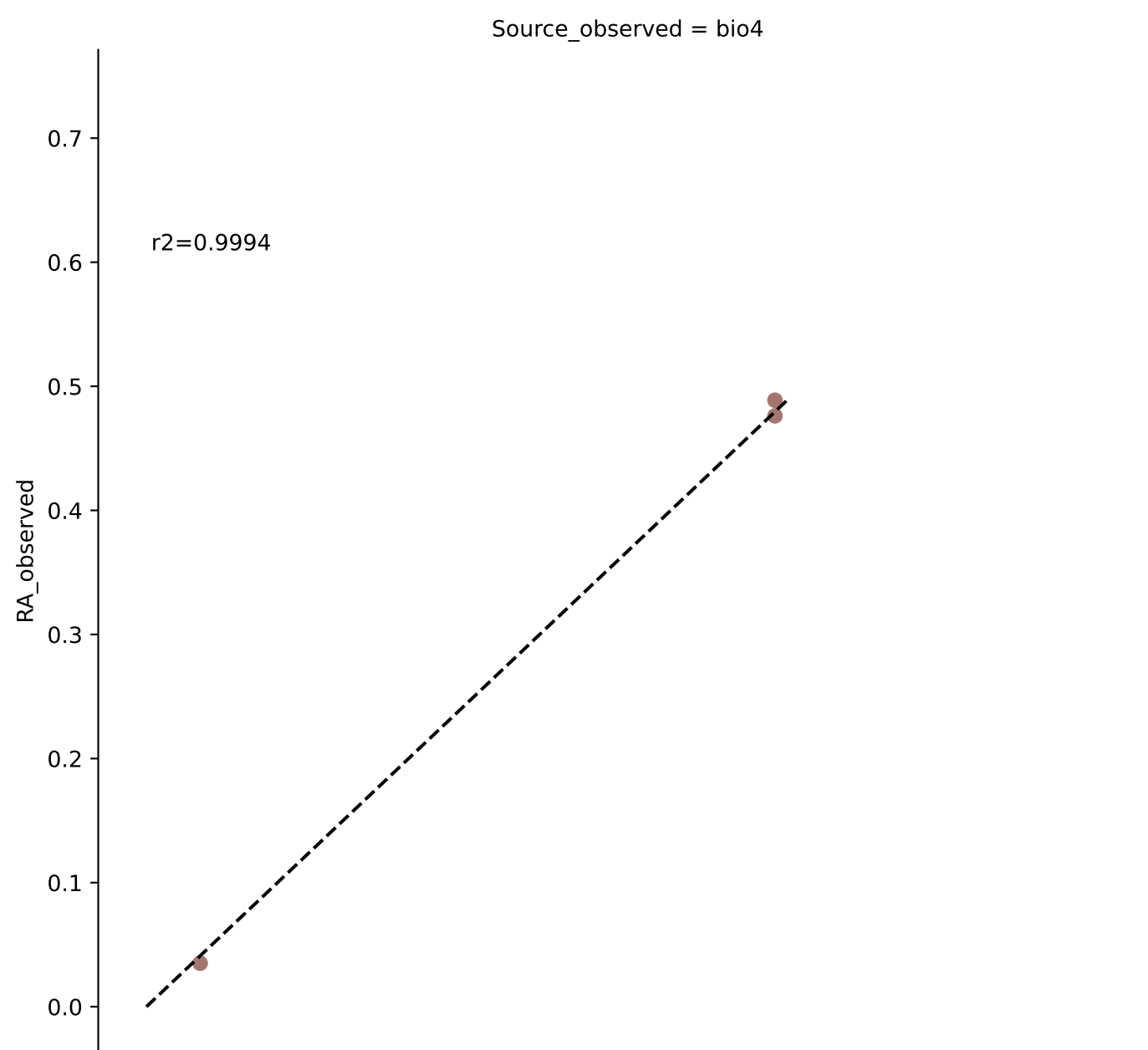


Bivariate Linear Regression for Sample EG in Experiment nist (Species at filter threshold 0.01)

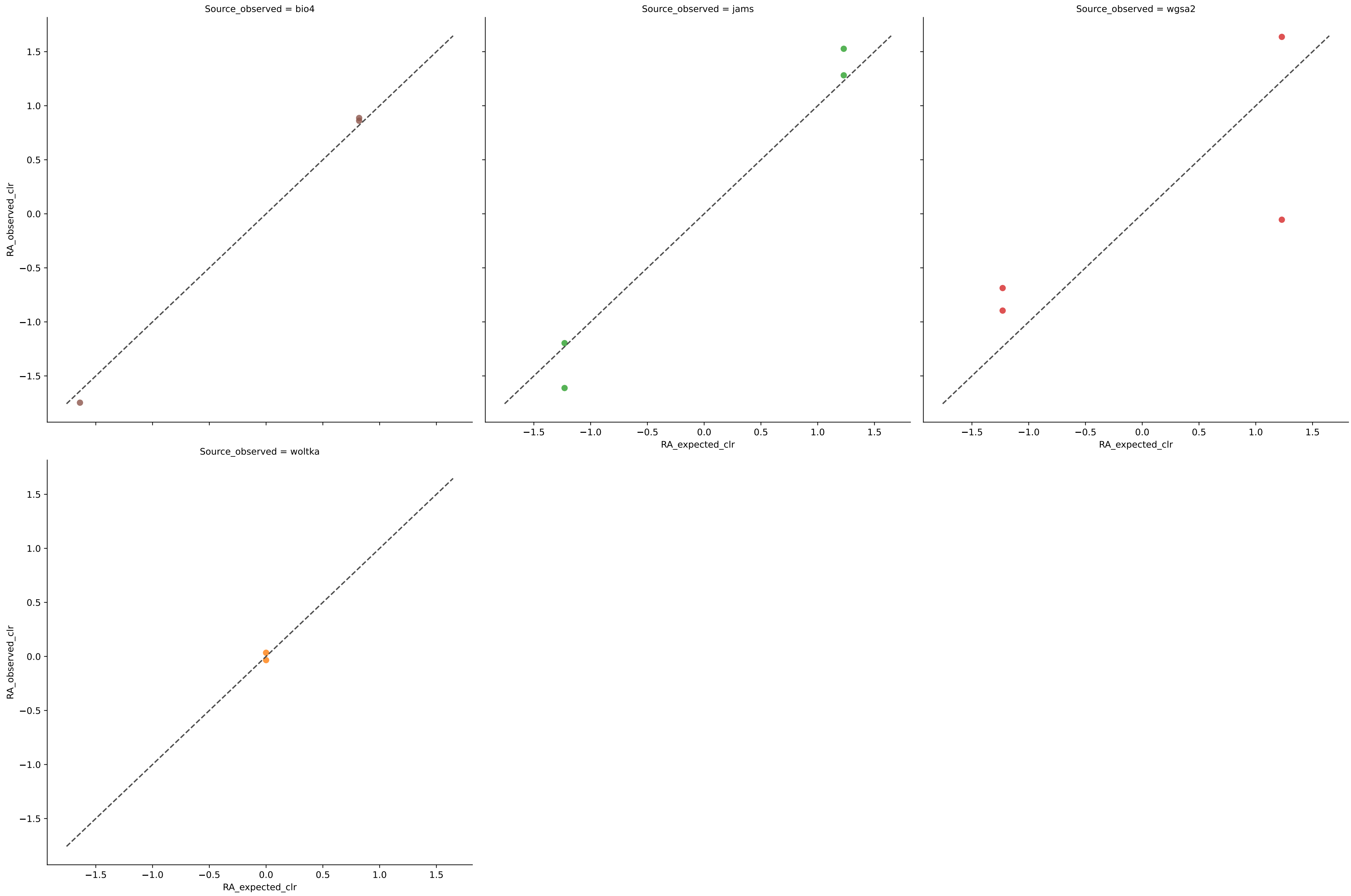


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	9	0.3670	0.0298	1.6749	0.8659	0.0361	100.0000	0.0000
jams	9	0.4711	0.0217	0.6955	0.9025	0.0286	100.0000	0.0000
wgsa2	8	0.3011	0.0485	1.1827	0.8059	0.0523	100.0000	0.0000
woltka	8	0.5676	0.0900	1.9558	0.6401	0.1052	100.0000	0.0000

Bivariate Linear Regression for Sample MIX-A in Experiment nist (Species at filter threshold 0.01)

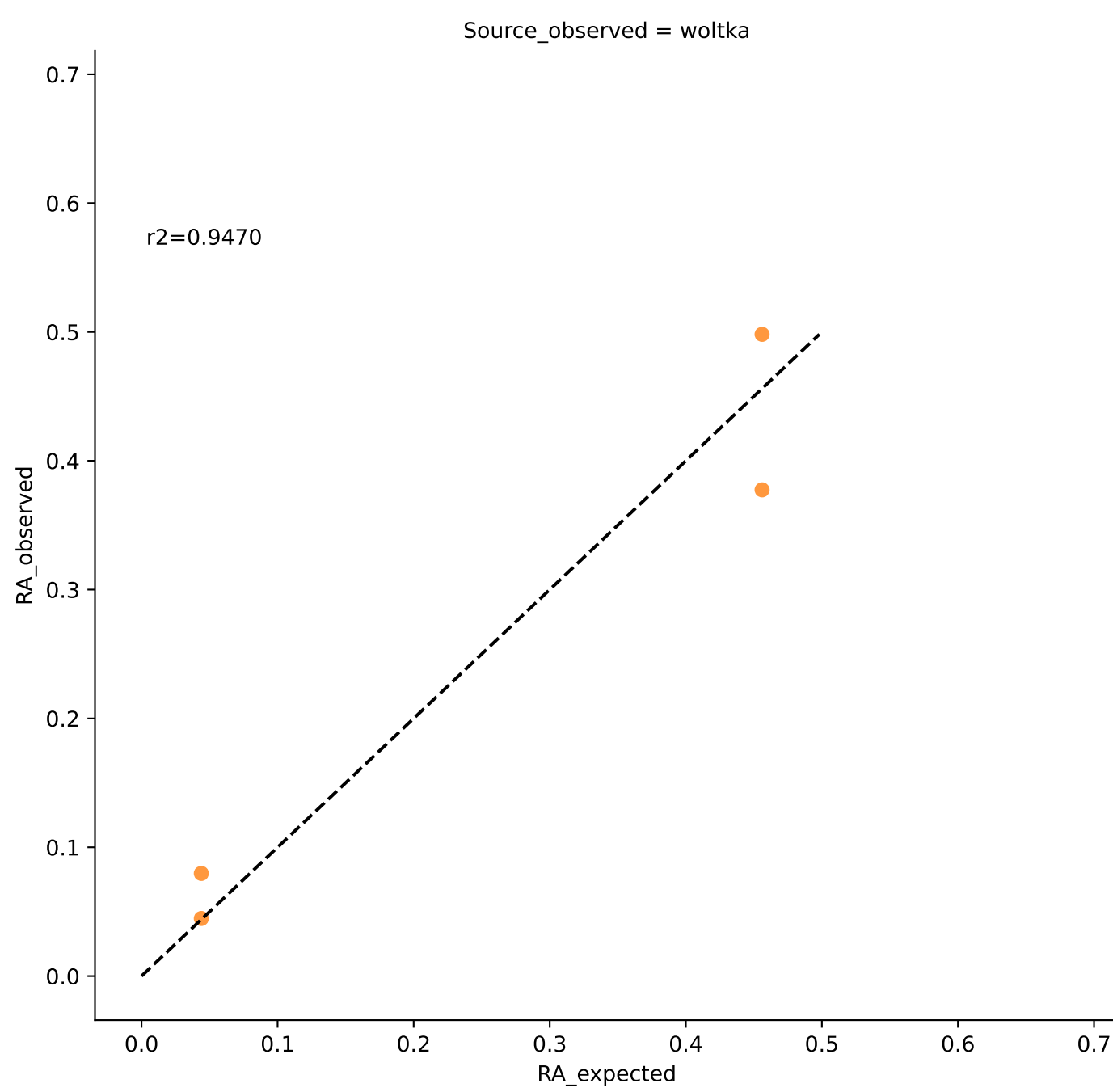
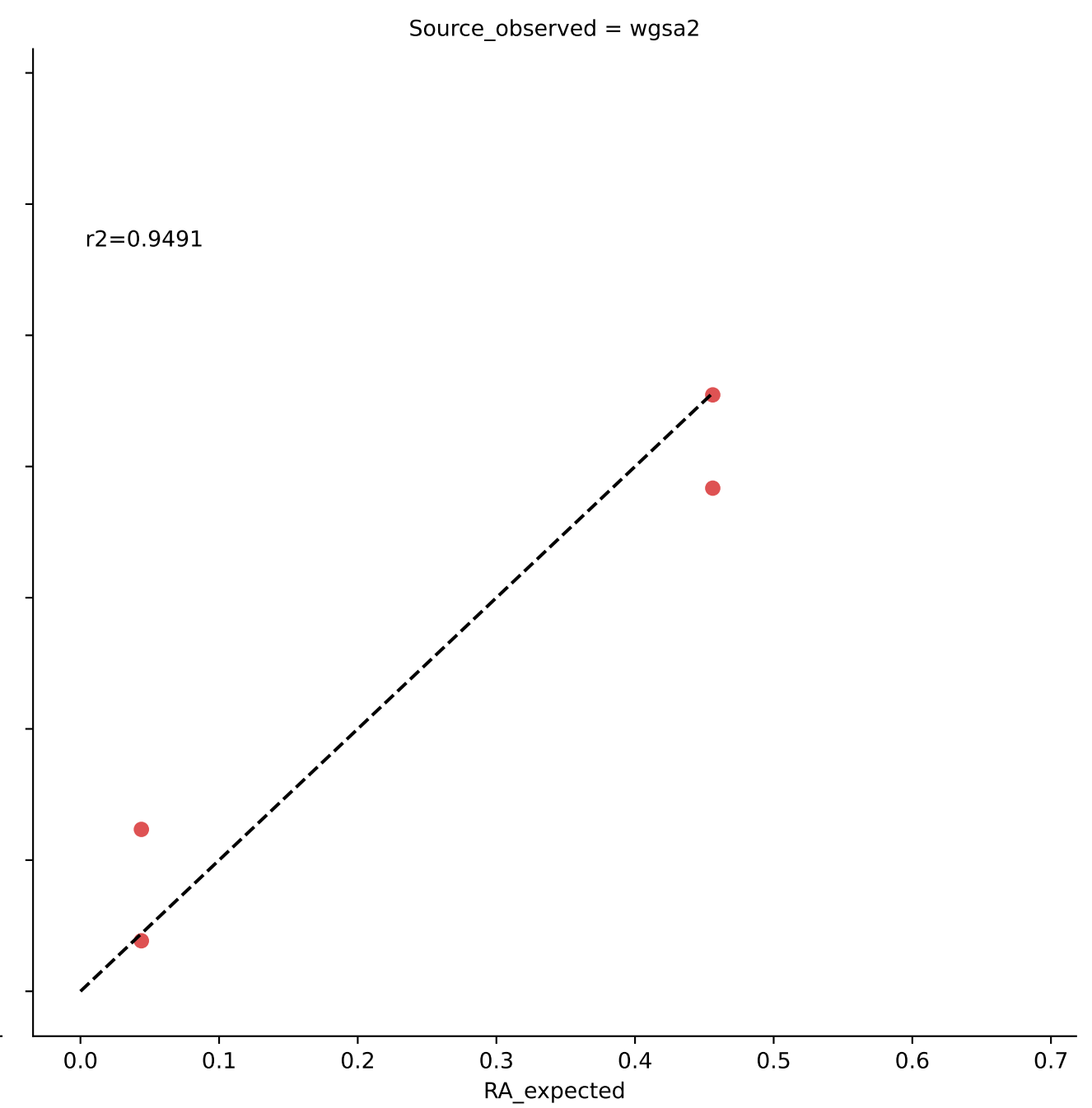
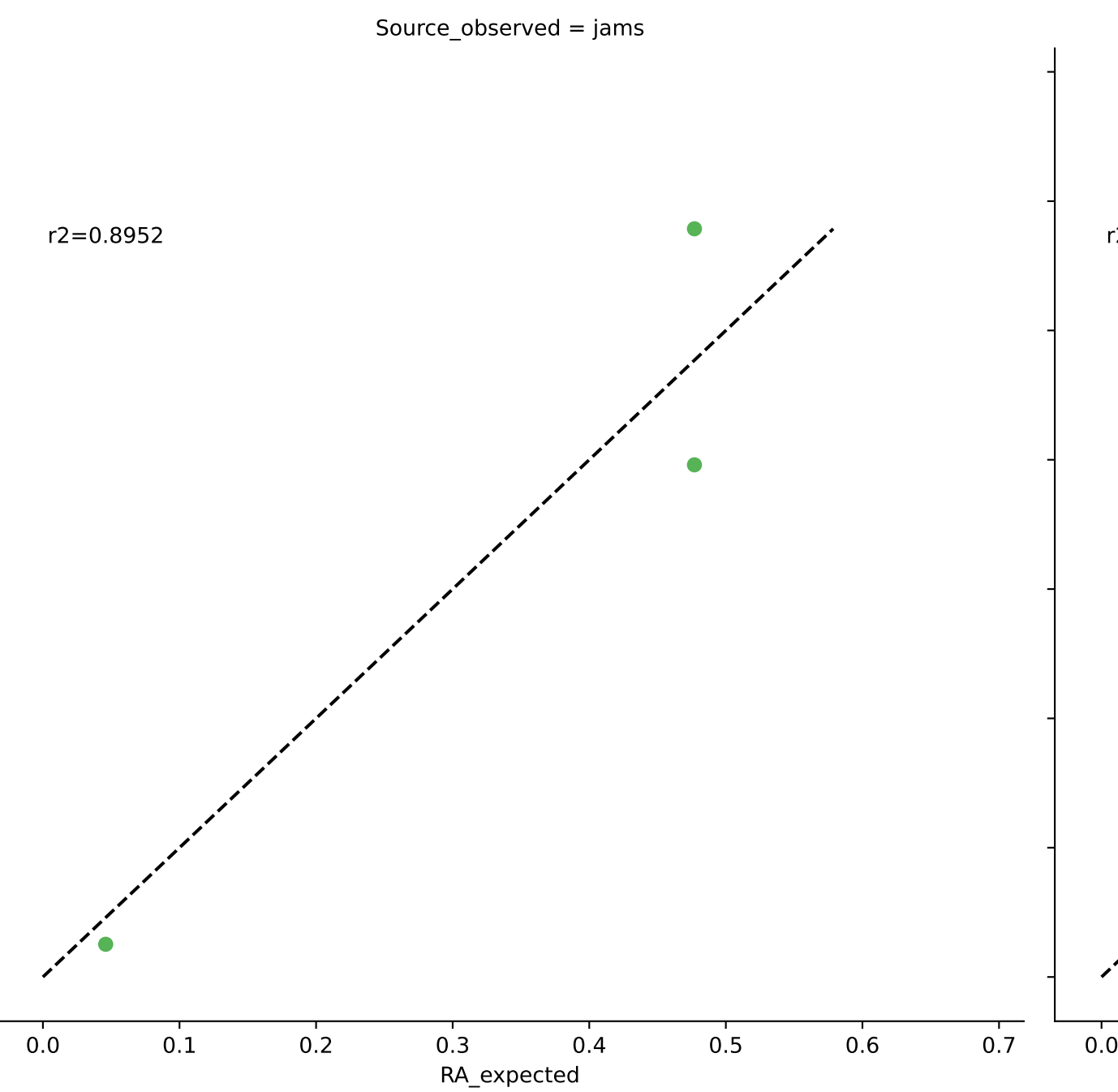
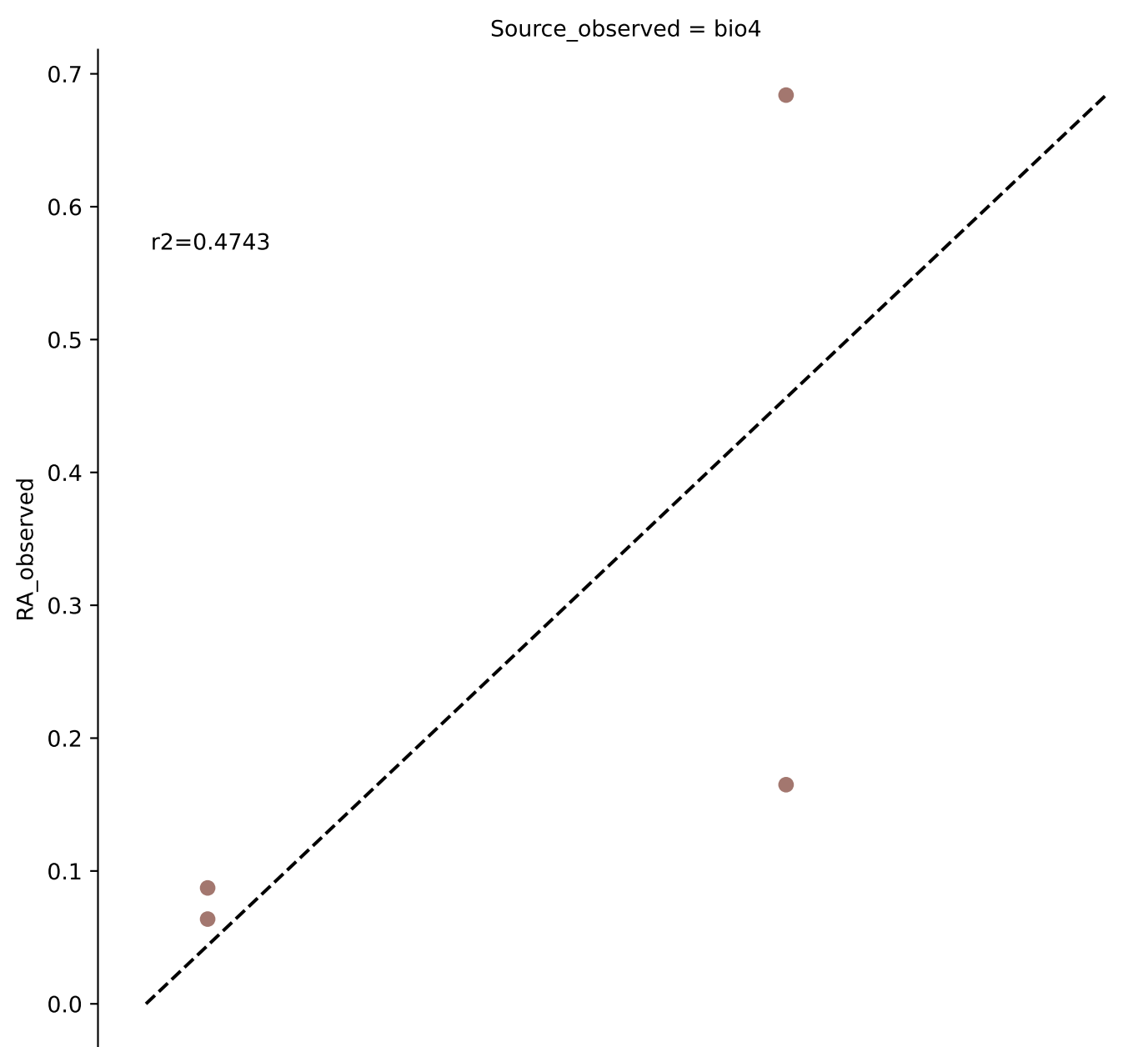


Bivariate Linear Regression for Sample MIX-A in Experiment nist (Species at filter threshold 0.01)

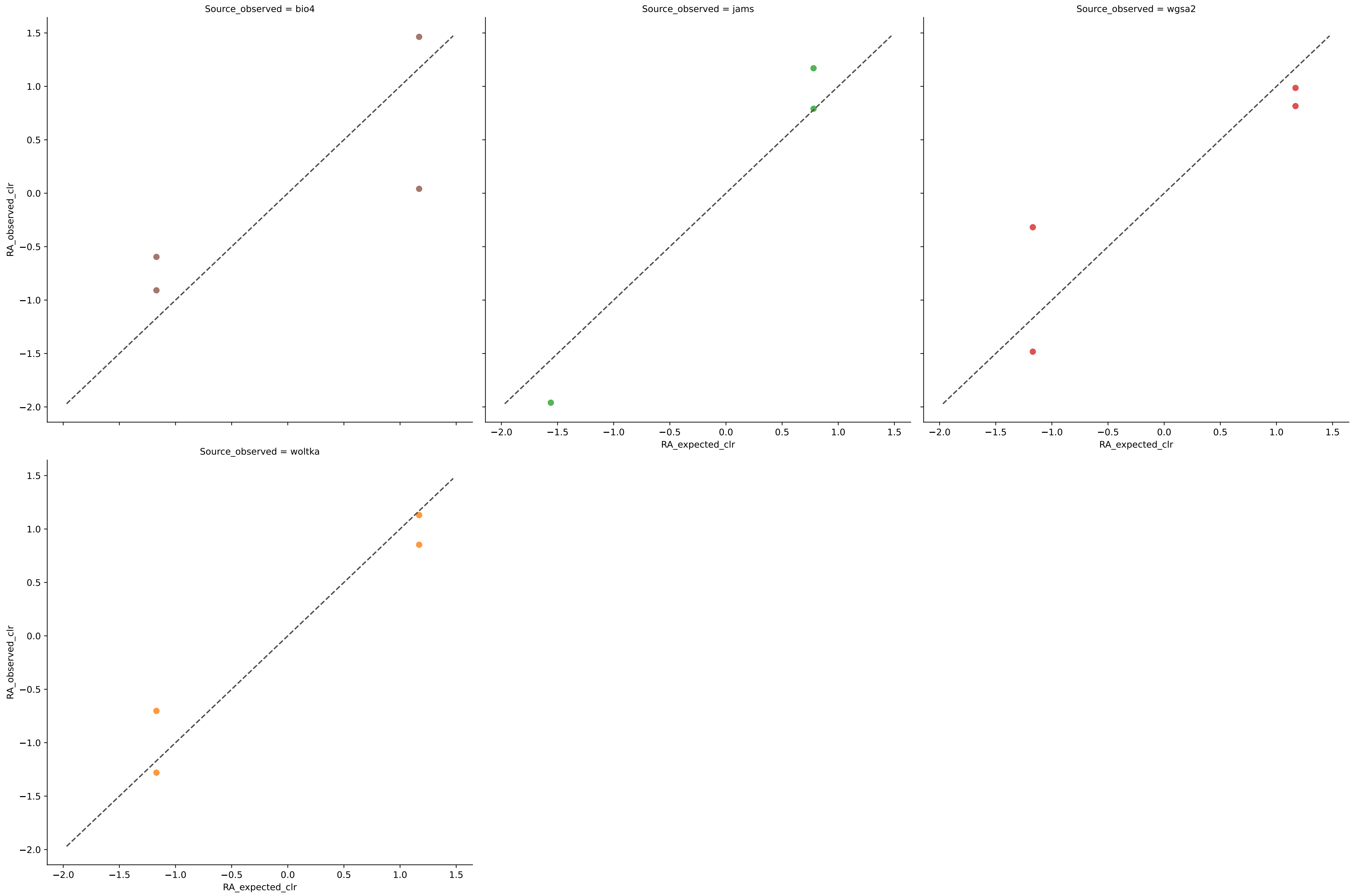


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	3	0.9994	0.0062	0.1338	0.9908	0.0066	100.0000	0.0000
jams	4	0.9669	0.0341	0.4885	0.9319	0.0422	100.0000	0.0000
wgsa2	4	0.4322	0.1627	1.4902	0.6747	0.2134	100.0000	0.0000
woltka	2	1.0000	0.0170	0.0482	0.9830	0.0170	100.0000	0.0000

Bivariate Linear Regression for Sample MIX-B in Experiment nist (Species at filter threshold 0.01)

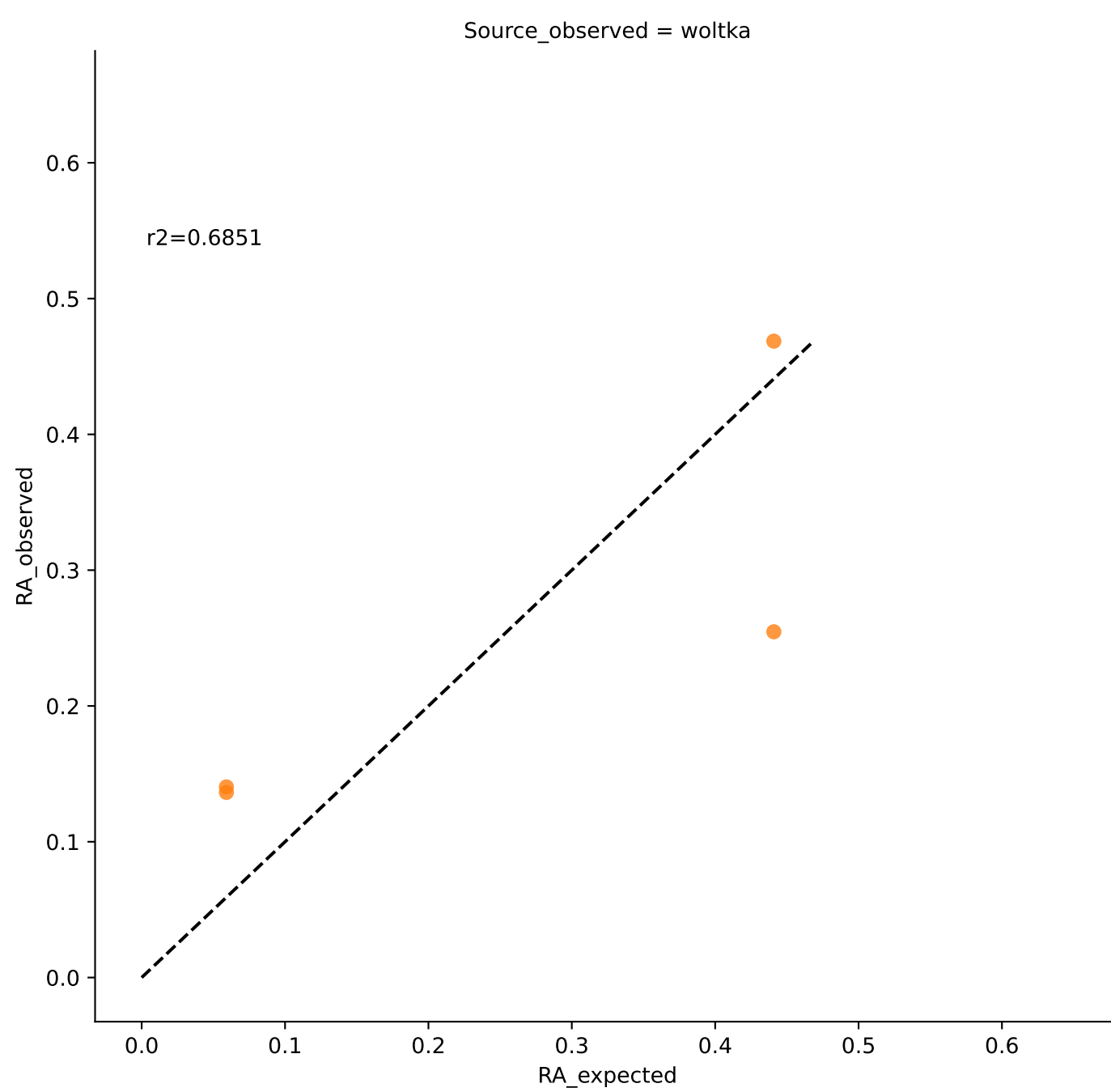
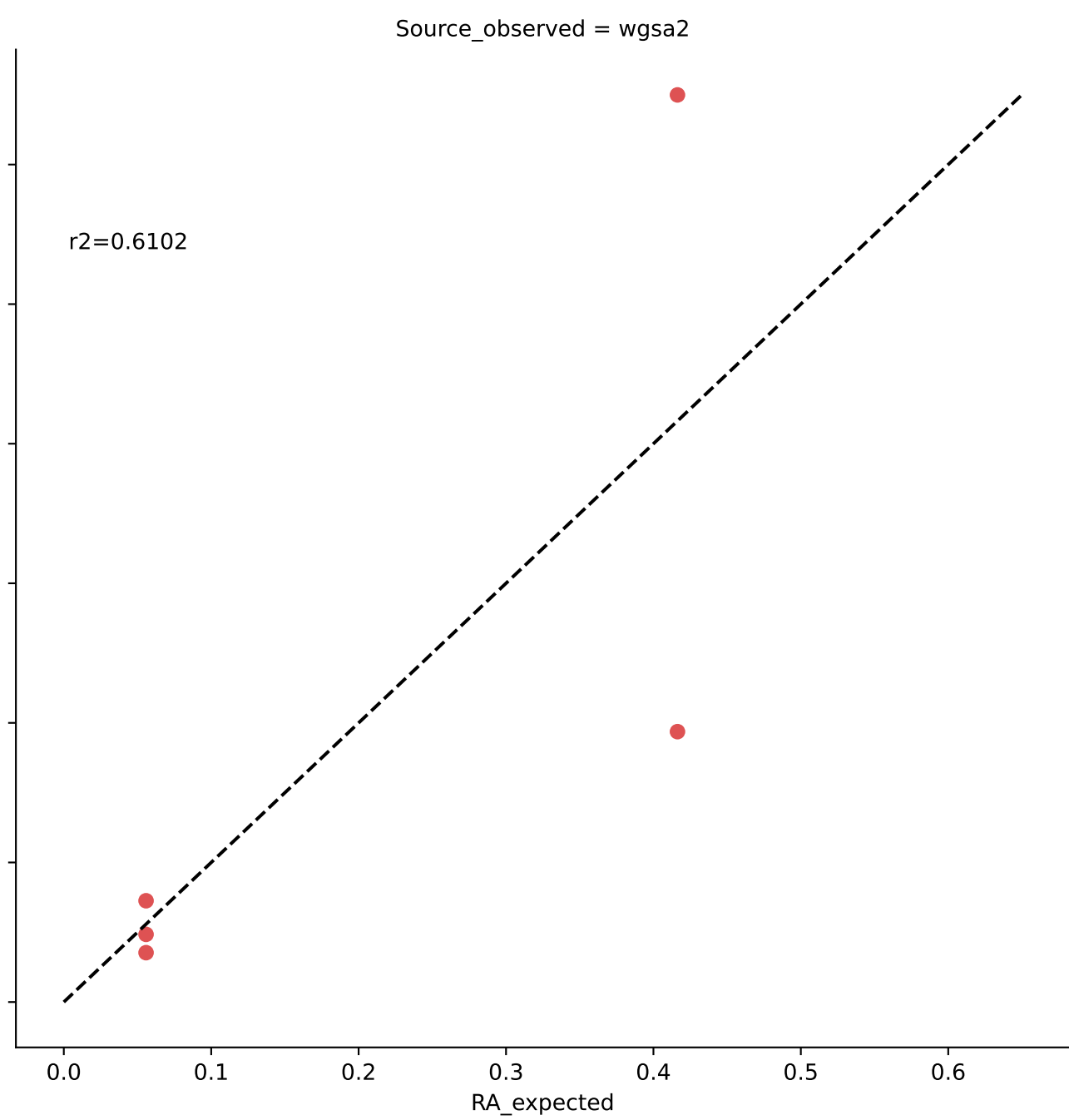
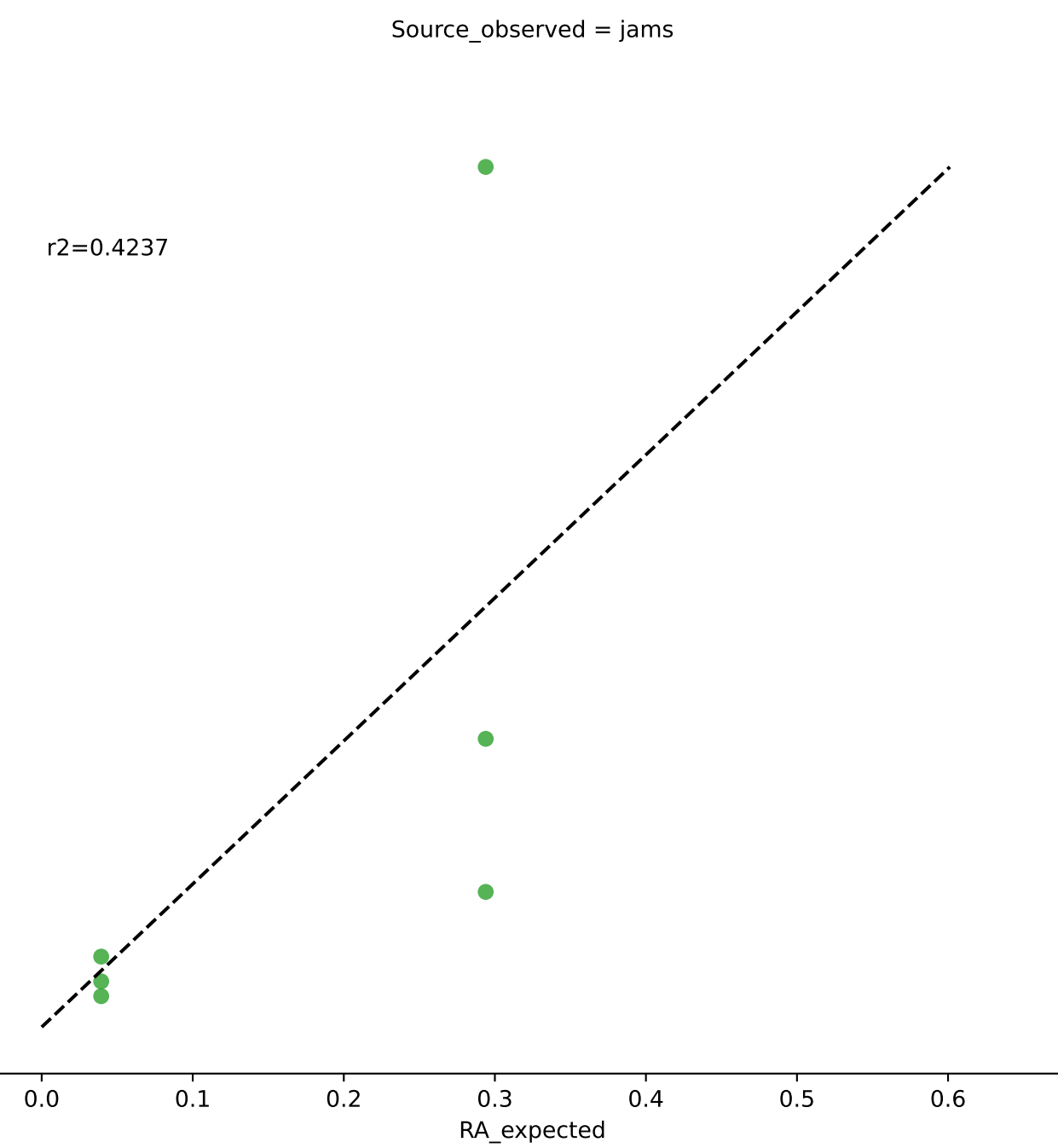
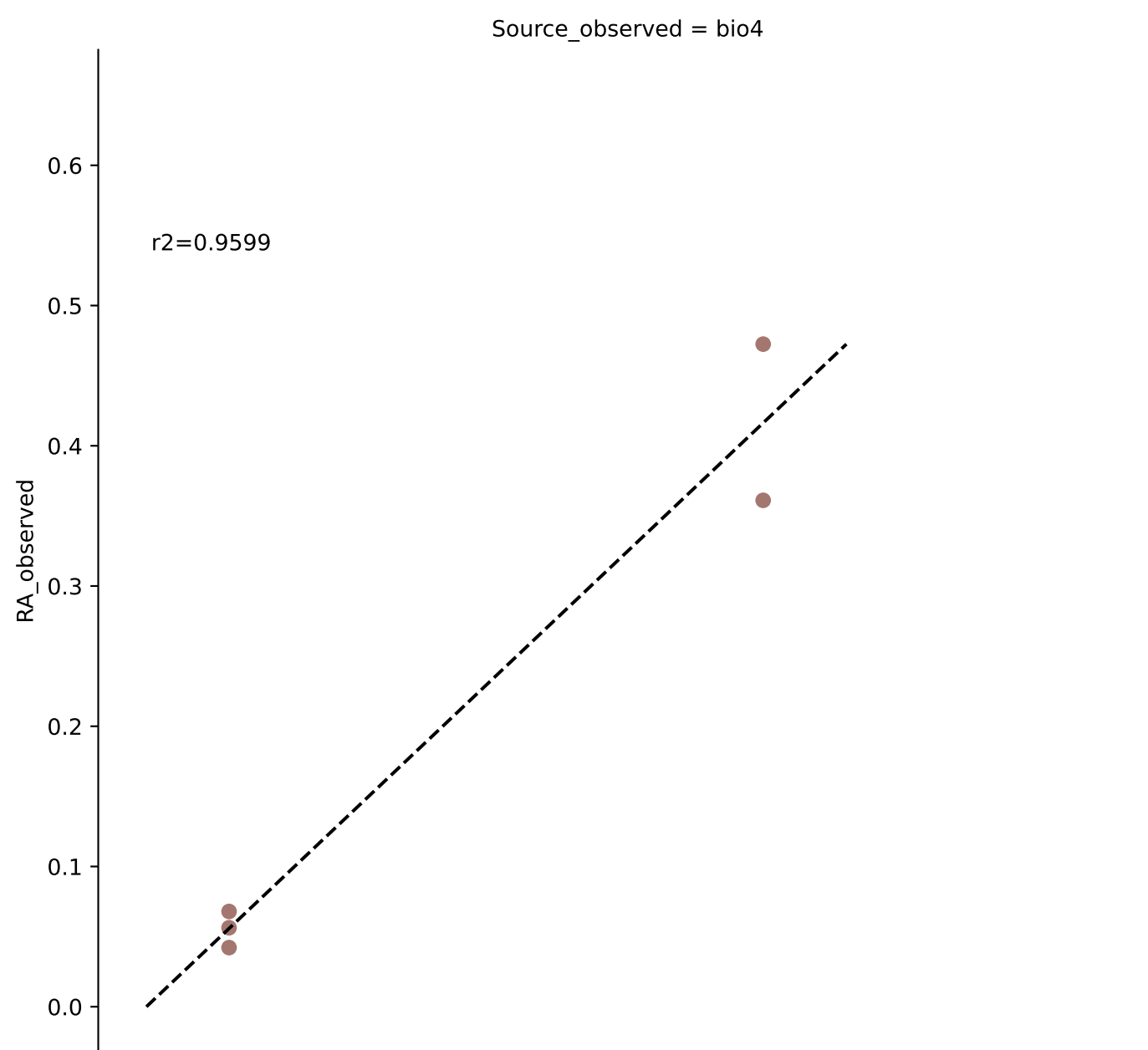


Bivariate Linear Regression for Sample MIX-B in Experiment nist (Species at filter threshold 0.01)

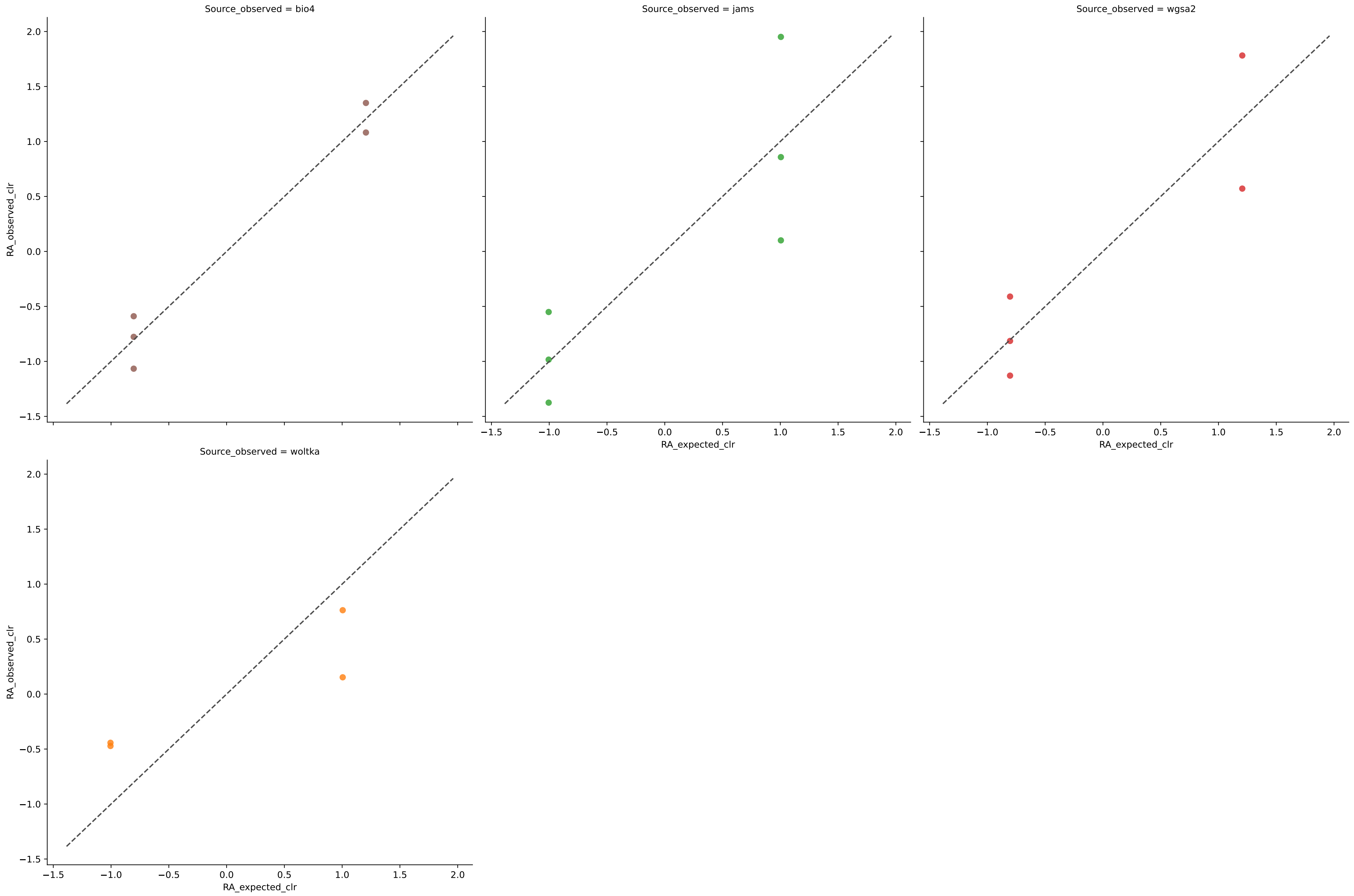


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	4	0.4743	0.1456	1.3260	0.7089	0.1864	100.0000	0.0000
jams	3	0.8952	0.0677	0.5589	0.8984	0.0759	100.0000	0.0000
wgsa2	4	0.9491	0.0397	0.9914	0.9205	0.0539	100.0000	0.0000
woltka	4	0.9470	0.0393	0.5771	0.9214	0.0481	100.0000	0.0000

Bivariate Linear Regression for Sample MIX-C in Experiment nist (Species at filter threshold 0.01)

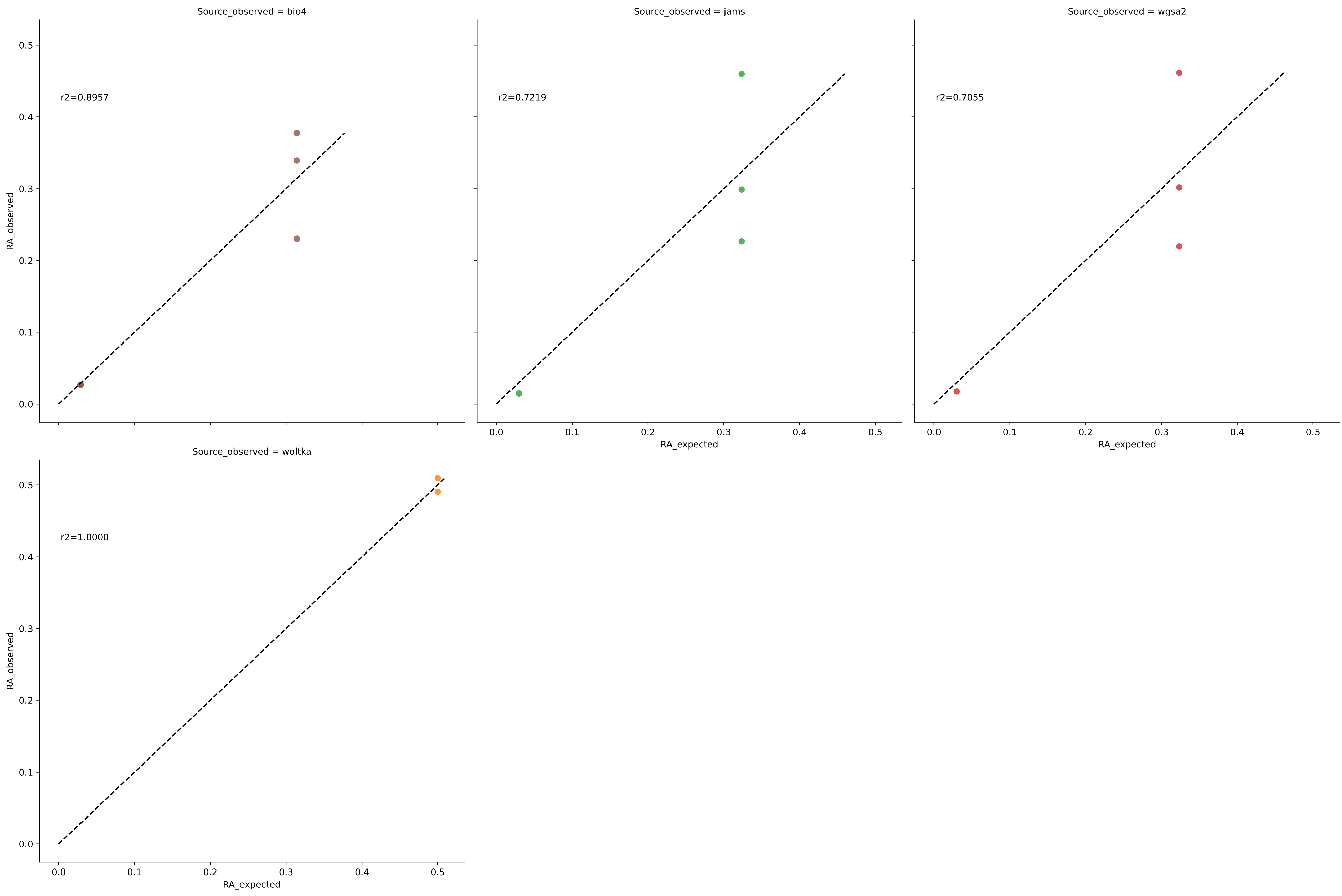


Bivariate Linear Regression for Sample MIX-C in Experiment nist (Species at filter threshold 0.01)

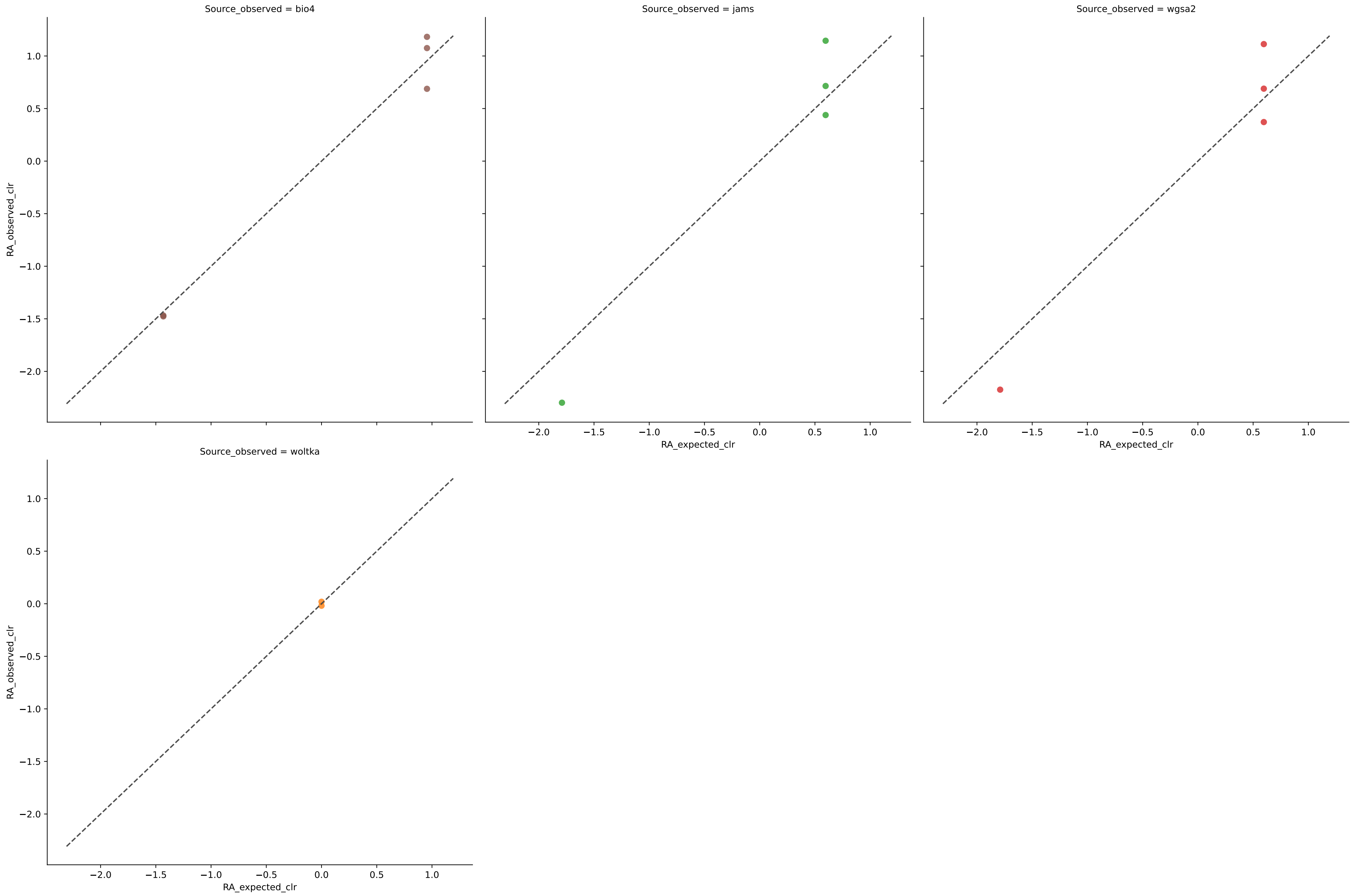


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	5	0.9599	0.0275	0.3898	0.9312	0.0362	100.0000	0.0000
jams	6	0.4237	0.1057	1.4416	0.6829	0.1545	100.0000	0.0000
wgsa2	5	0.6102	0.1001	0.9975	0.7497	0.1448	100.0000	0.0000
woltka	4	0.6851	0.0931	1.1768	0.8137	0.1096	100.0000	0.0000

Bivariate Linear Regression for Sample MIX-D in Experiment nist (Species at filter threshold 0.01)

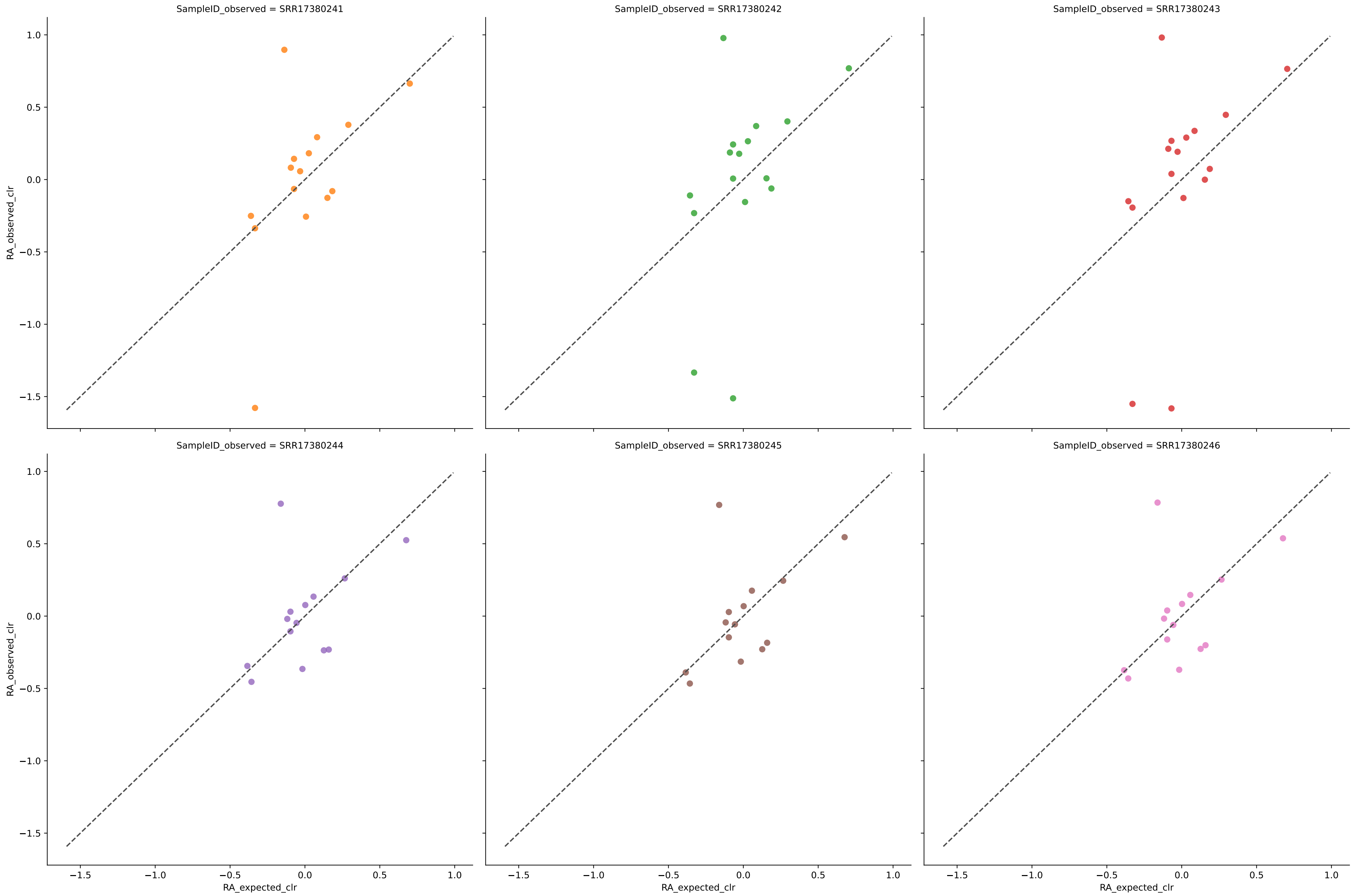


Bivariate Linear Regression for Sample MIX-D in Experiment nist (Species at filter threshold 0.01)



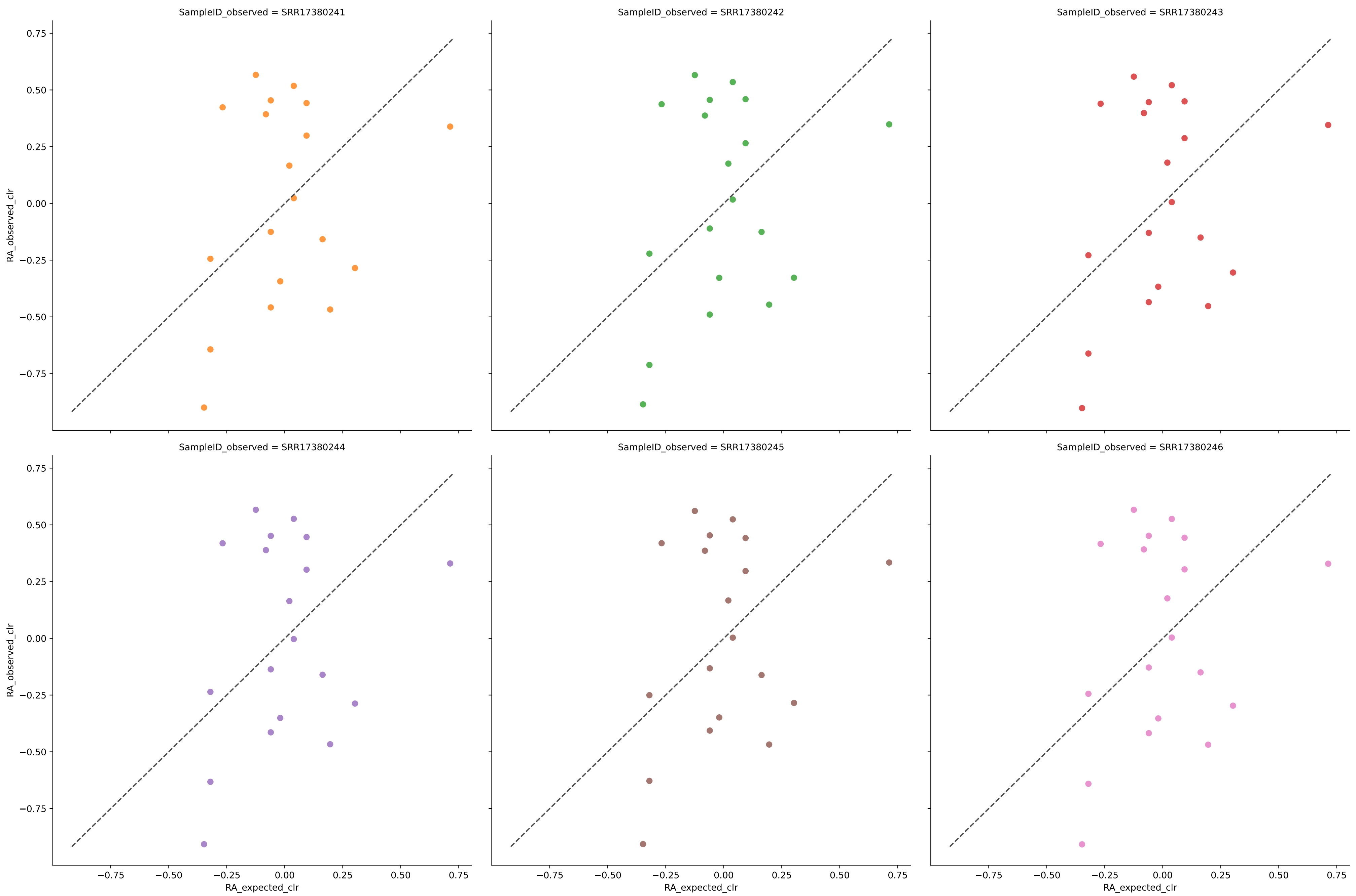
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
bio4	5	0.8957	0.0354	0.3753	0.9115	0.0484	100.0000	0.0000
jams	4	0.7219	0.0681	0.7732	0.8637	0.0848	100.0000	0.0000
wgsa2	4	0.7055	0.0689	0.6880	0.8622	0.0871	100.0000	0.0000
woltka	2	1.0000	0.0095	0.0268	0.9905	0.0095	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using bio4 in Experiment tourlousse with filter 0.01



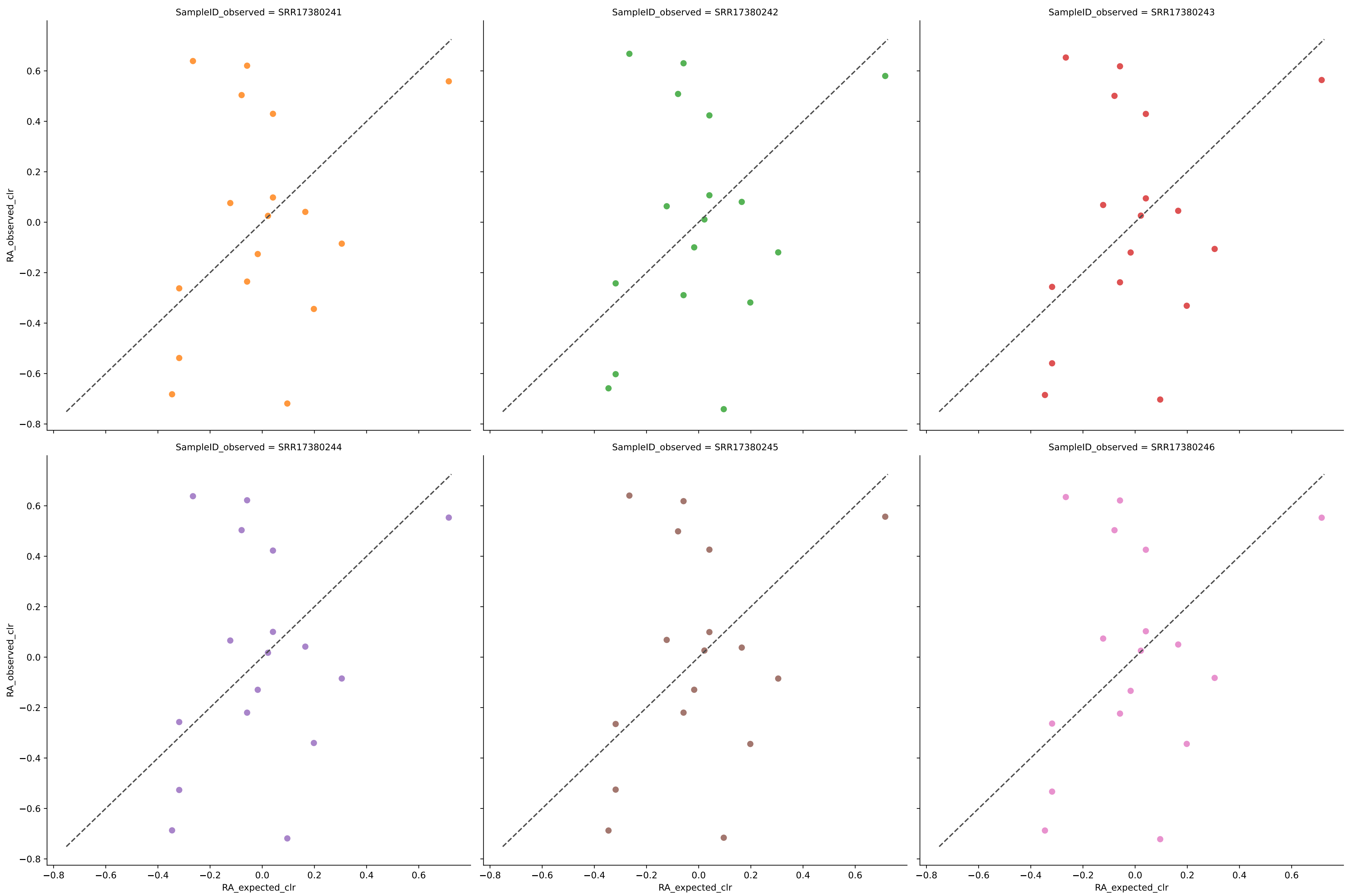
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR17380241	15	0.2457	0.0027	1.7353	0.8797	0.0045	100.0000	0.0000
SRR17380242	16	0.2317	0.0031	2.2095	0.8583	0.0047	100.0000	0.0000
SRR17380243	16	0.2397	0.0030	2.3586	0.8628	0.0047	100.0000	0.0000
SRR17380244	14	0.1678	0.0025	1.1651	0.8935	0.0044	100.0000	0.0000
SRR17380245	14	0.1959	0.0024	1.1265	0.8976	0.0043	100.0000	0.0000
SRR17380246	14	0.1753	0.0025	1.1606	0.8941	0.0044	100.0000	0.0000
Average	15	0.2093	0.0027	1.6260	0.8810	0.0045	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using jams in Experiment tourlousse with filter 0.01



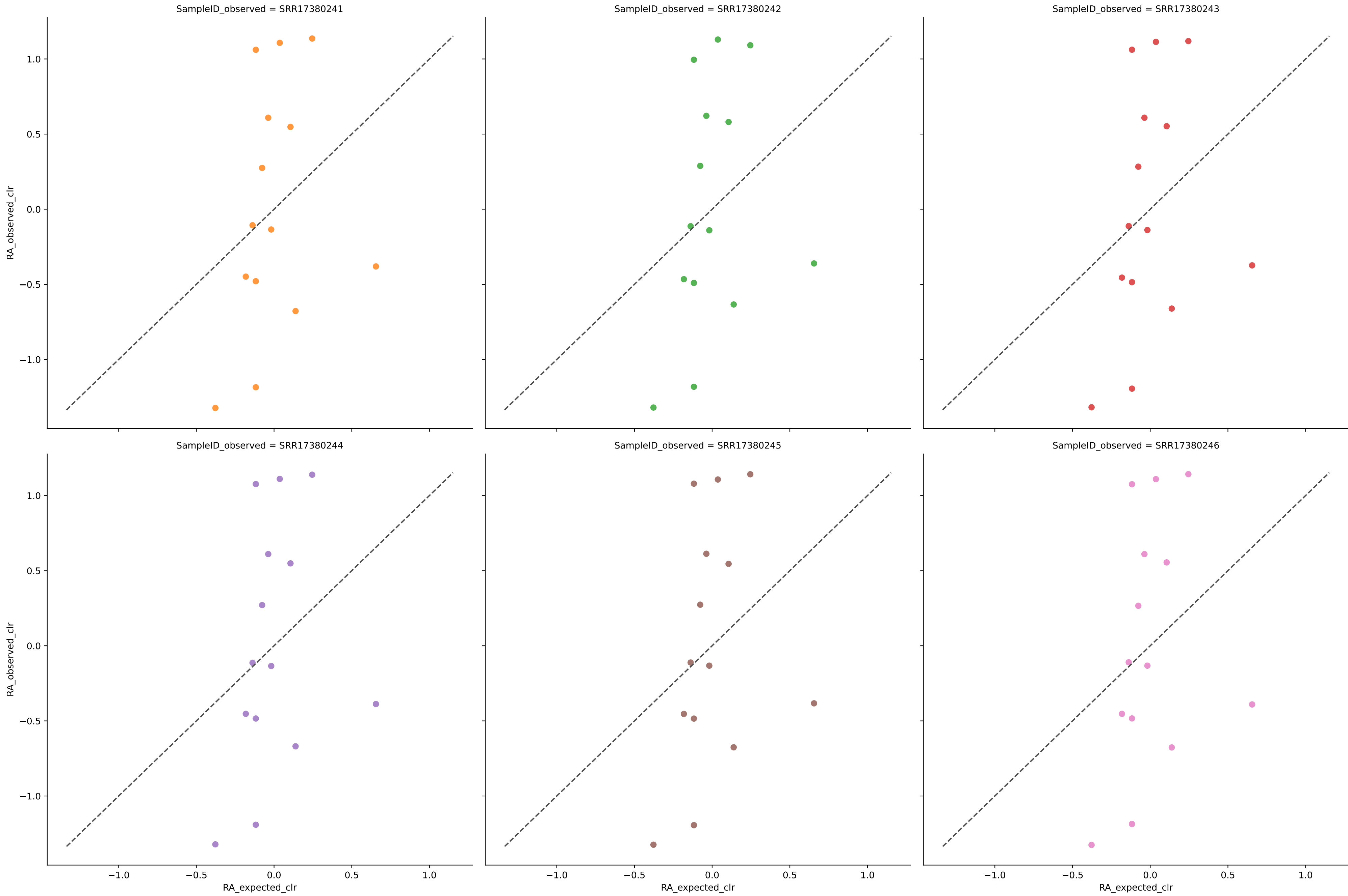
	Diversity	R ²	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR17380241	19	0.0294	0.0033	1.8893	0.8104	0.0039	100.0000	0.0000
SRR17380242	19	0.0298	0.0033	1.9110	0.8093	0.0039	100.0000	0.0000
SRR17380243	19	0.0296	0.0033	1.8964	0.8091	0.0039	100.0000	0.0000
SRR17380244	19	0.0276	0.0033	1.8859	0.8093	0.0039	100.0000	0.0000
SRR17380245	19	0.0291	0.0033	1.8776	0.8108	0.0039	100.0000	0.0000
SRR17380246	19	0.0276	0.0033	1.8902	0.8094	0.0039	100.0000	0.0000
Average	19	0.0289	0.0033	1.8917	0.8097	0.0039	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using wgsa in Experiment tourlousse with filter 0.01



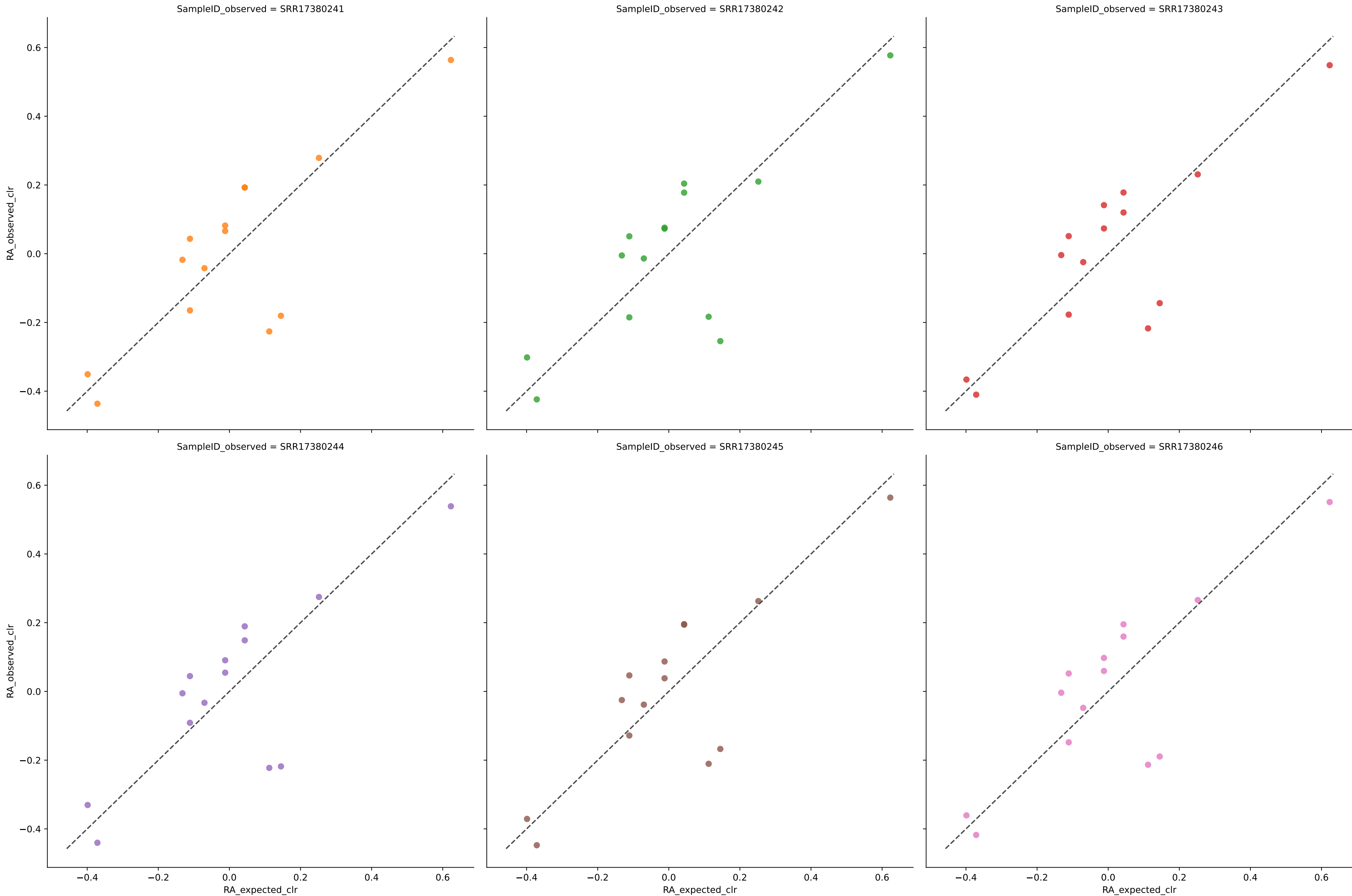
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR17380241	17	0.0700	0.0034	1.7806	0.8270	0.0044	100.0000	0.0000
SRR17380242	17	0.0716	0.0034	1.8147	0.8245	0.0044	100.0000	0.0000
SRR17380243	17	0.0699	0.0034	1.7812	0.8269	0.0044	100.0000	0.0000
SRR17380244	17	0.0684	0.0034	1.7755	0.8273	0.0044	100.0000	0.0000
SRR17380245	17	0.0698	0.0034	1.7745	0.8279	0.0044	100.0000	0.0000
SRR17380246	17	0.0693	0.0034	1.7778	0.8274	0.0044	100.0000	0.0000
Average	17	0.0698	0.0034	1.7840	0.8268	0.0044	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using woltka in Experiment tourlousse with filter 0.01



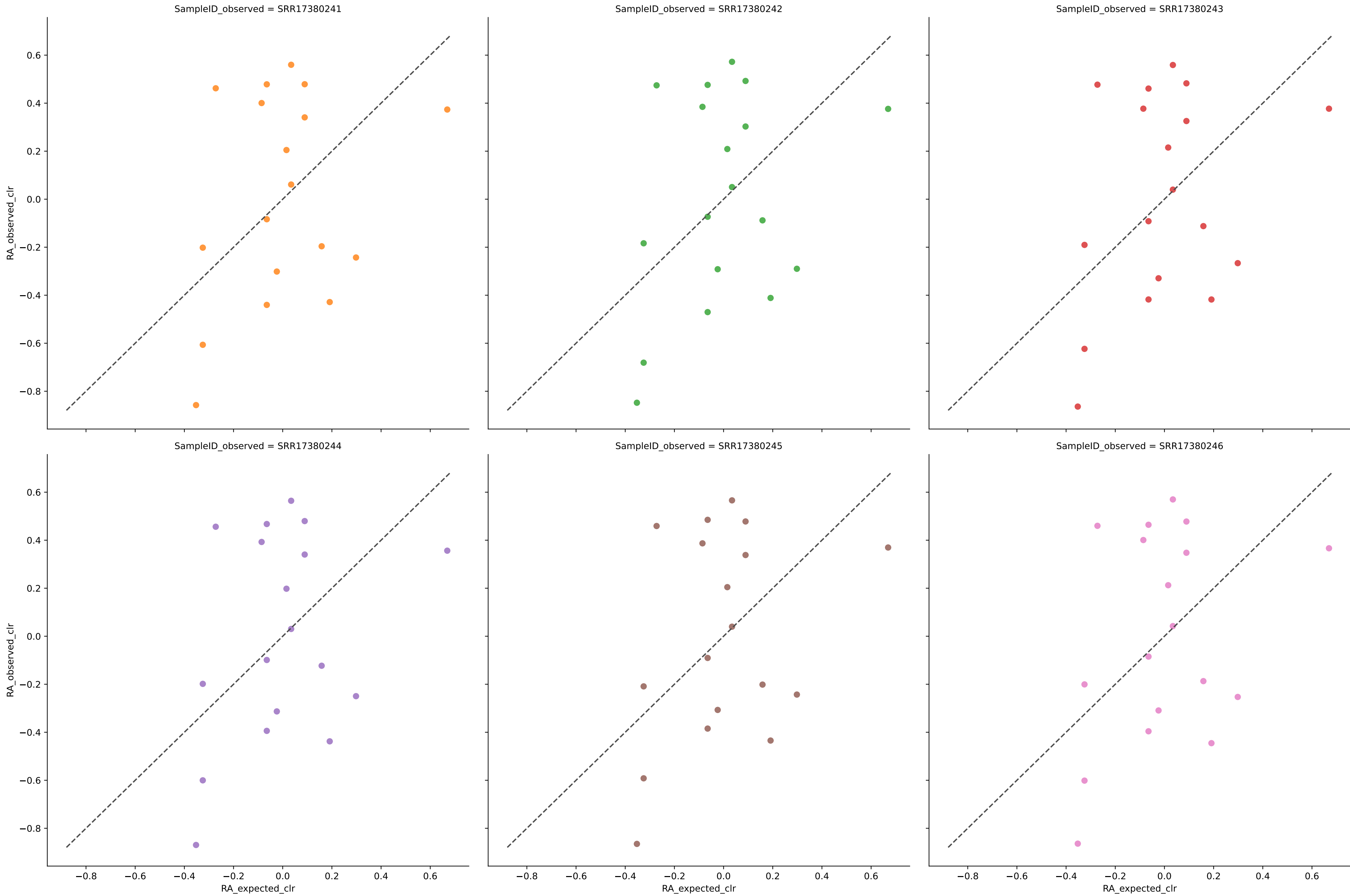
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR17380241	14	0.0101	0.0075	2.8383	0.6862	0.0091	100.0000	0.0000
SRR17380242	14	0.0114	0.0074	2.7974	0.6893	0.0088	100.0000	0.0000
SRR17380243	14	0.0099	0.0075	2.8335	0.6862	0.0090	100.0000	0.0000
SRR17380244	14	0.0097	0.0075	2.8495	0.6848	0.0091	100.0000	0.0000
SRR17380245	14	0.0098	0.0075	2.8526	0.6847	0.0091	100.0000	0.0000
SRR17380246	14	0.0097	0.0075	2.8525	0.6846	0.0091	100.0000	0.0000
Average	14	0.0101	0.0075	2.8373	0.6860	0.0090	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using bio4 in Experiment tourlousse with filter 0.01



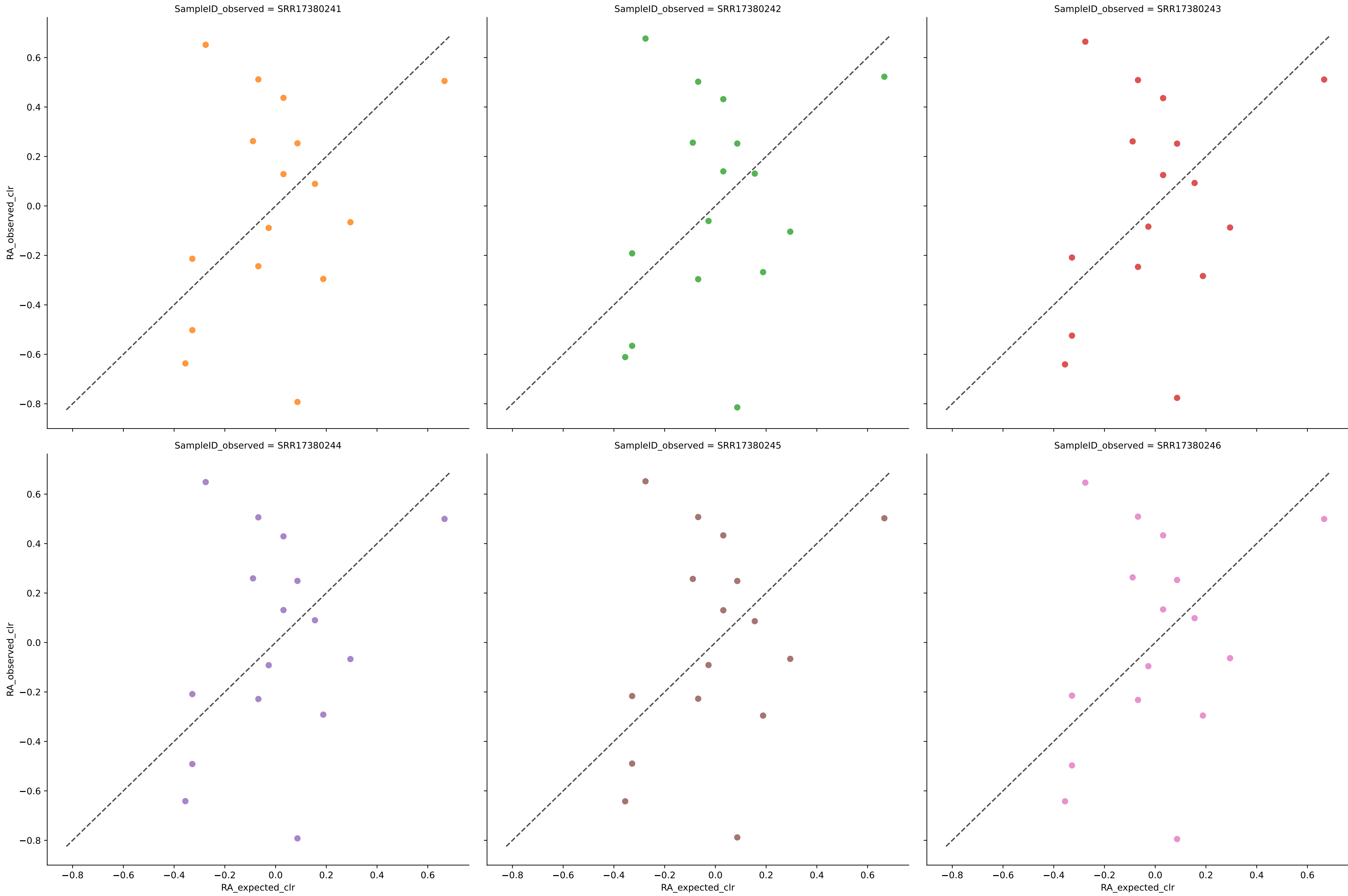
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR17380241	14	0.7188	0.0014	0.5758	0.9413	0.0018	100.0000	0.0000
SRR17380242	14	0.6939	0.0015	0.6104	0.9375	0.0018	100.0000	0.0000
SRR17380243	14	0.7251	0.0014	0.5518	0.9418	0.0017	100.0000	0.0000
SRR17380244	14	0.6958	0.0014	0.5919	0.9399	0.0018	100.0000	0.0000
SRR17380245	14	0.7324	0.0013	0.5549	0.9449	0.0017	100.0000	0.0000
SRR17380246	14	0.7145	0.0014	0.5704	0.9423	0.0018	100.0000	0.0000
Average	14	0.7134	0.0014	0.5759	0.9413	0.0018	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using jams in Experiment tourlousse with filter 0.01



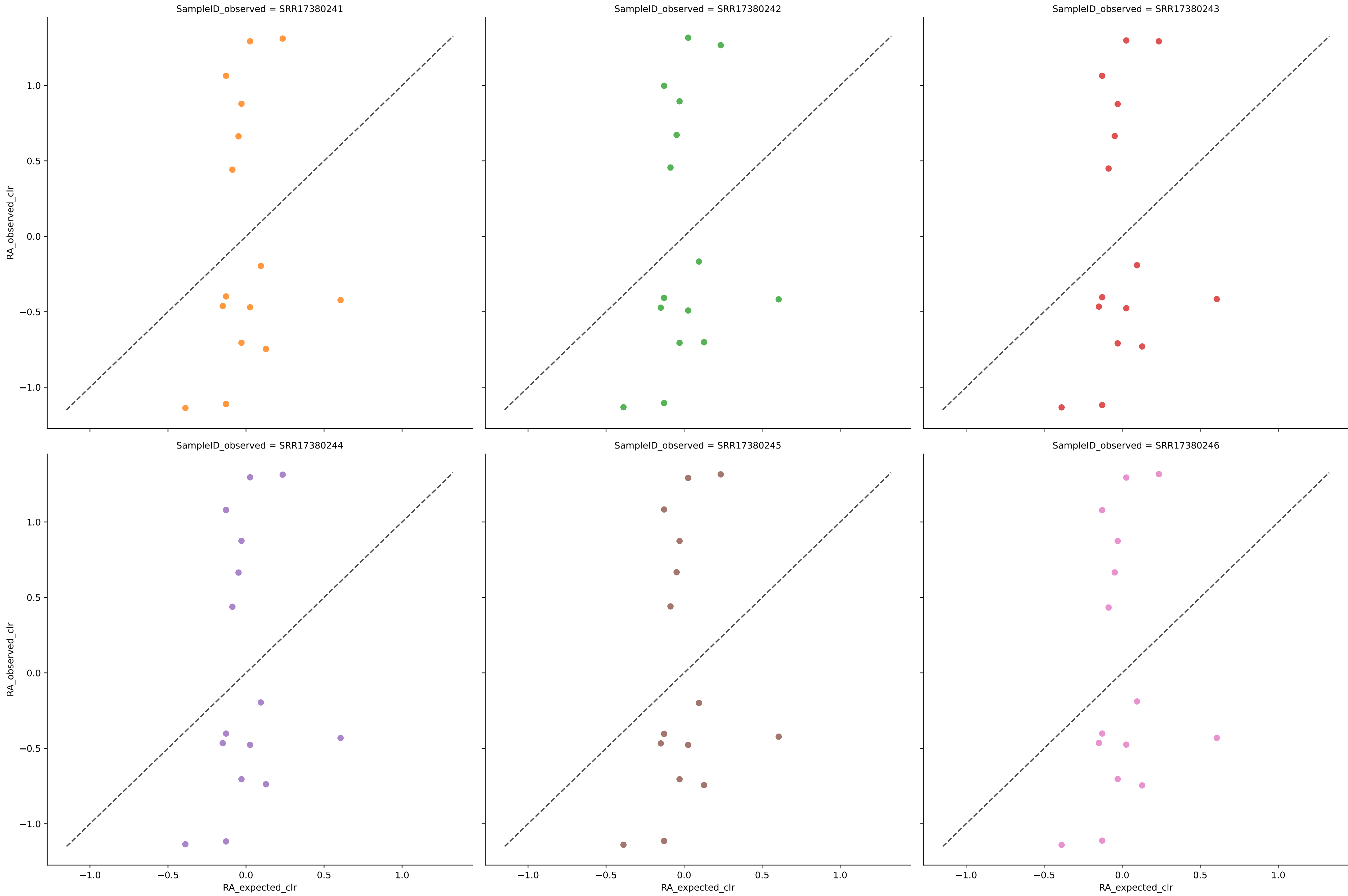
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR17380241	18	0.0539	0.0033	1.7483	0.8195	0.0039	100.0000	0.0000
SRR17380242	18	0.0554	0.0033	1.7580	0.8201	0.0039	100.0000	0.0000
SRR17380243	18	0.0570	0.0033	1.7359	0.8207	0.0038	100.0000	0.0000
SRR17380244	18	0.0525	0.0033	1.7313	0.8202	0.0039	100.0000	0.0000
SRR17380245	18	0.0529	0.0033	1.7396	0.8196	0.0039	100.0000	0.0000
SRR17380246	18	0.0515	0.0034	1.7496	0.8187	0.0039	100.0000	0.0000
Average	18	0.0539	0.0033	1.7438	0.8198	0.0039	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using wgsa in Experiment tourlousse with filter 0.01



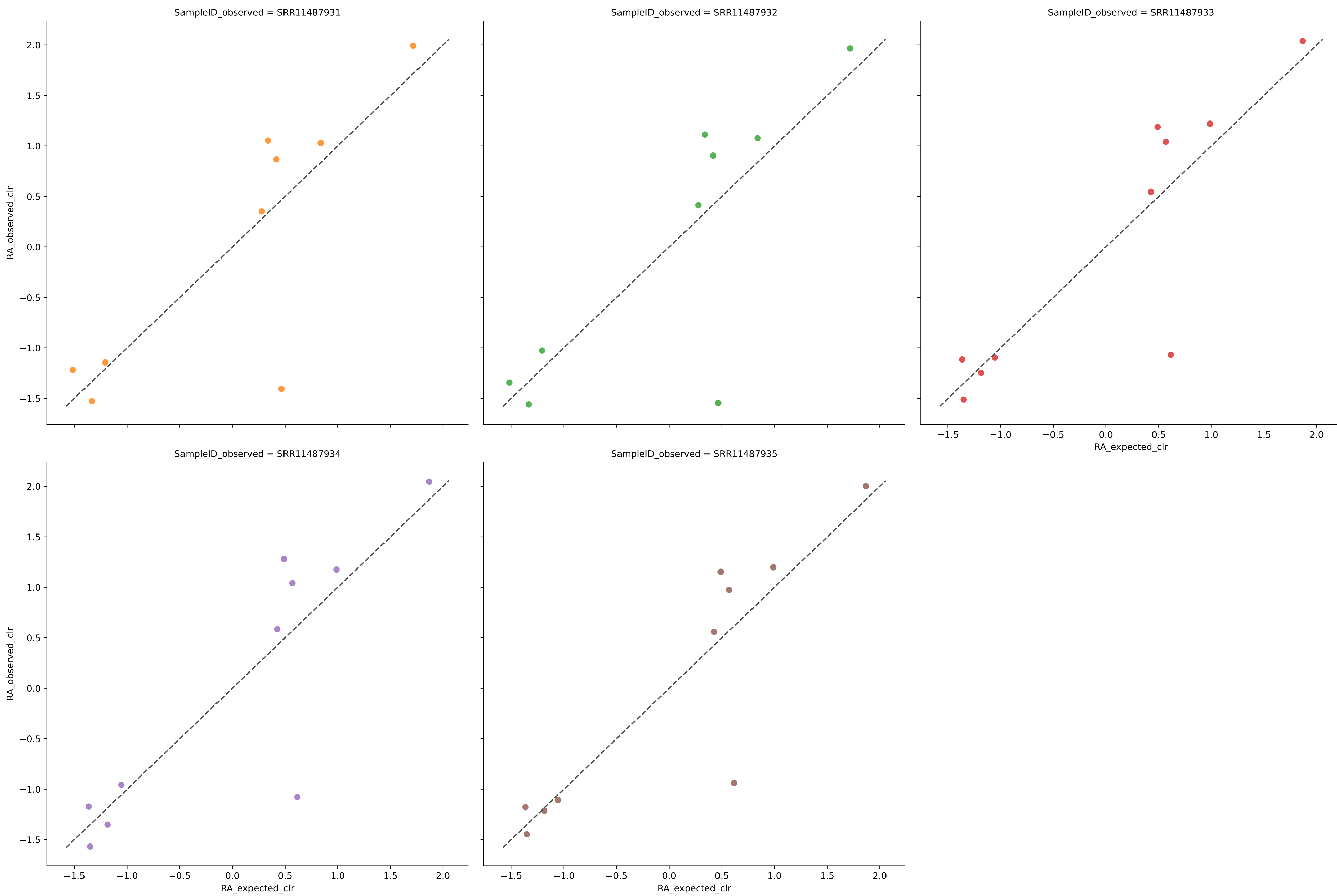
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR17380241	16	0.0716	0.0034	1.6867	0.8352	0.0044	100.0000	0.0000
SRR17380242	16	0.0730	0.0034	1.7159	0.8342	0.0044	100.0000	0.0000
SRR17380243	16	0.0711	0.0034	1.6876	0.8348	0.0044	100.0000	0.0000
SRR17380244	16	0.0707	0.0034	1.6785	0.8362	0.0044	100.0000	0.0000
SRR17380245	16	0.0710	0.0034	1.6791	0.8362	0.0044	100.0000	0.0000
SRR17380246	16	0.0714	0.0034	1.6827	0.8357	0.0044	100.0000	0.0000
Average	16	0.0715	0.0034	1.6884	0.8354	0.0044	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using woltka in Experiment tourlousse with filter 0.01



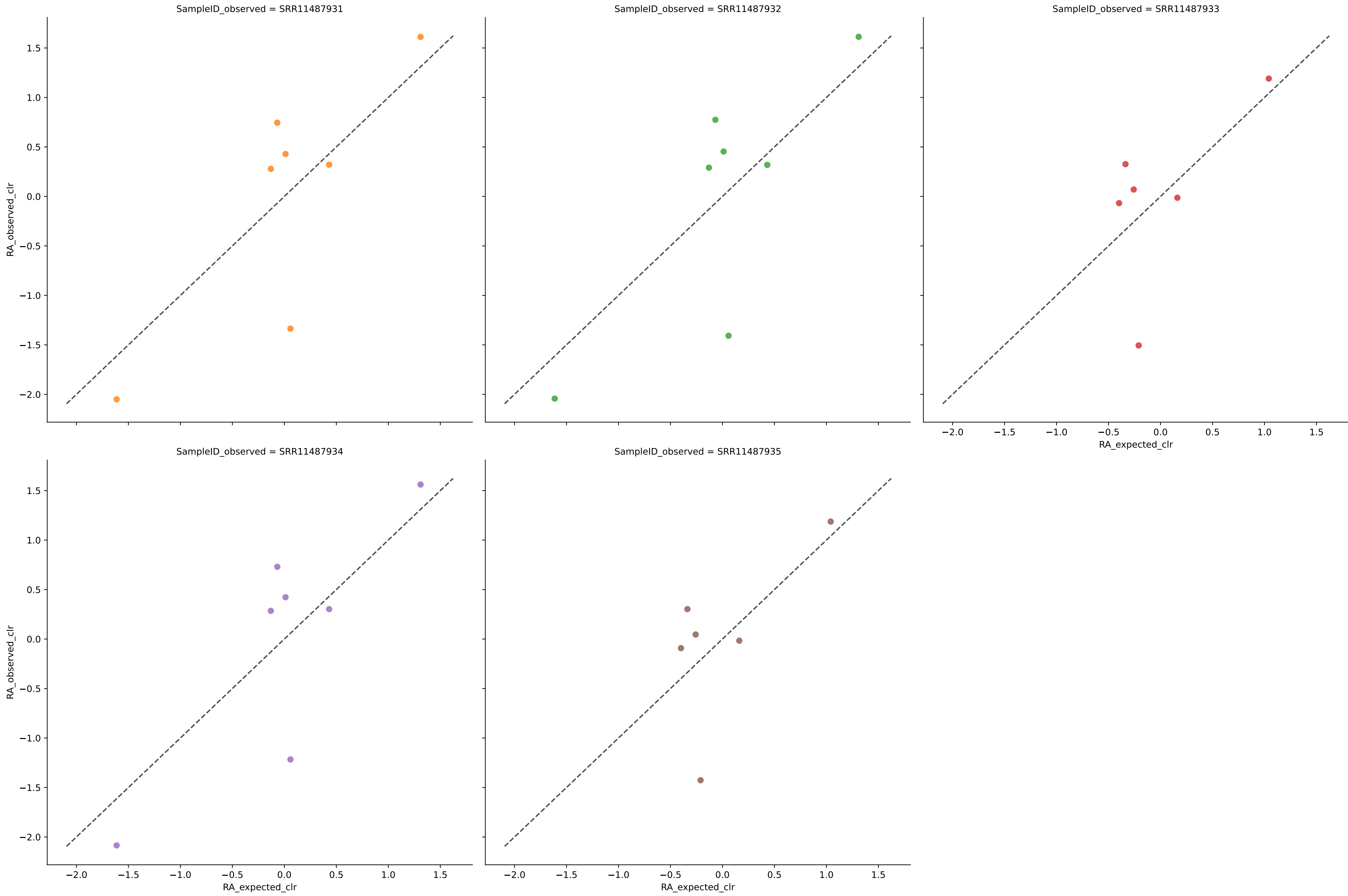
	Diversity	R ²	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR17380241	15	0.0043	0.0083	3.1779	0.6280	0.0094	100.0000	0.0000
SRR17380242	15	0.0042	0.0082	3.1438	0.6316	0.0093	100.0000	0.0000
SRR17380243	15	0.0040	0.0083	3.1722	0.6285	0.0094	100.0000	0.0000
SRR17380244	15	0.0040	0.0083	3.1884	0.6270	0.0095	100.0000	0.0000
SRR17380245	15	0.0042	0.0083	3.1886	0.6267	0.0095	100.0000	0.0000
SRR17380246	15	0.0042	0.0083	3.1875	0.6272	0.0095	100.0000	0.0000
Average	15	0.0041	0.0083	3.1764	0.6282	0.0094	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using bio4 in Experiment Amos hilo with filter 0.01



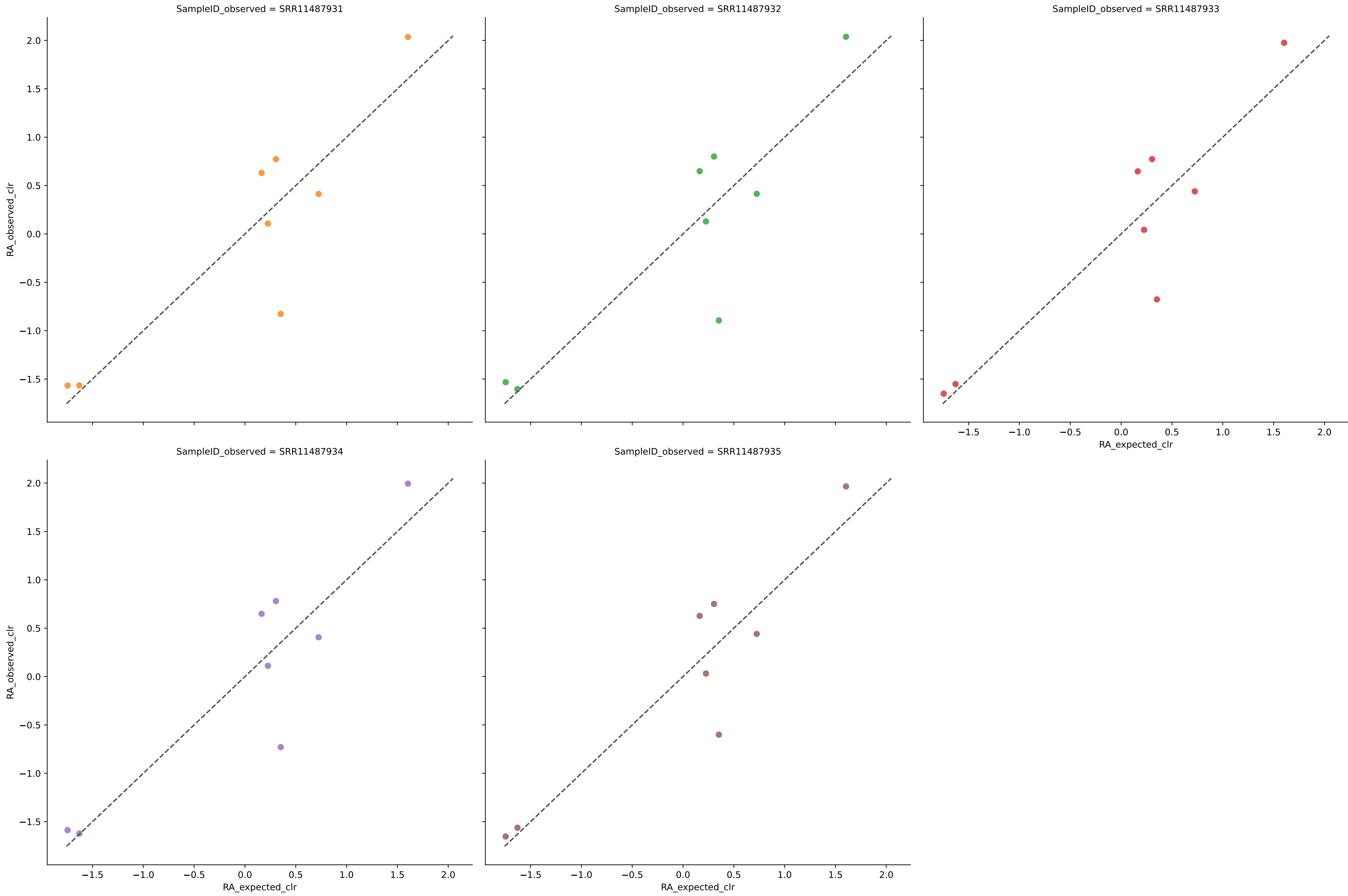
	Diversity	R ²	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487931	9	0.8903	0.0053	2.1140	0.8800	0.0081	100.0000	0.0000
SRR11487932	9	0.8730	0.0051	2.2639	0.8849	0.0084	100.0000	0.0000
SRR11487933	10	0.8851	0.0045	1.9349	0.8889	0.0075	100.0000	0.0000
SRR11487934	10	0.8704	0.0045	1.9826	0.8873	0.0080	100.0000	0.0000
SRR11487935	10	0.8906	0.0042	1.7737	0.8946	0.0073	100.0000	0.0000
Average	10	0.8819	0.0047	2.0138	0.8871	0.0078	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using jams in Experiment Amos hilo with filter 0.01



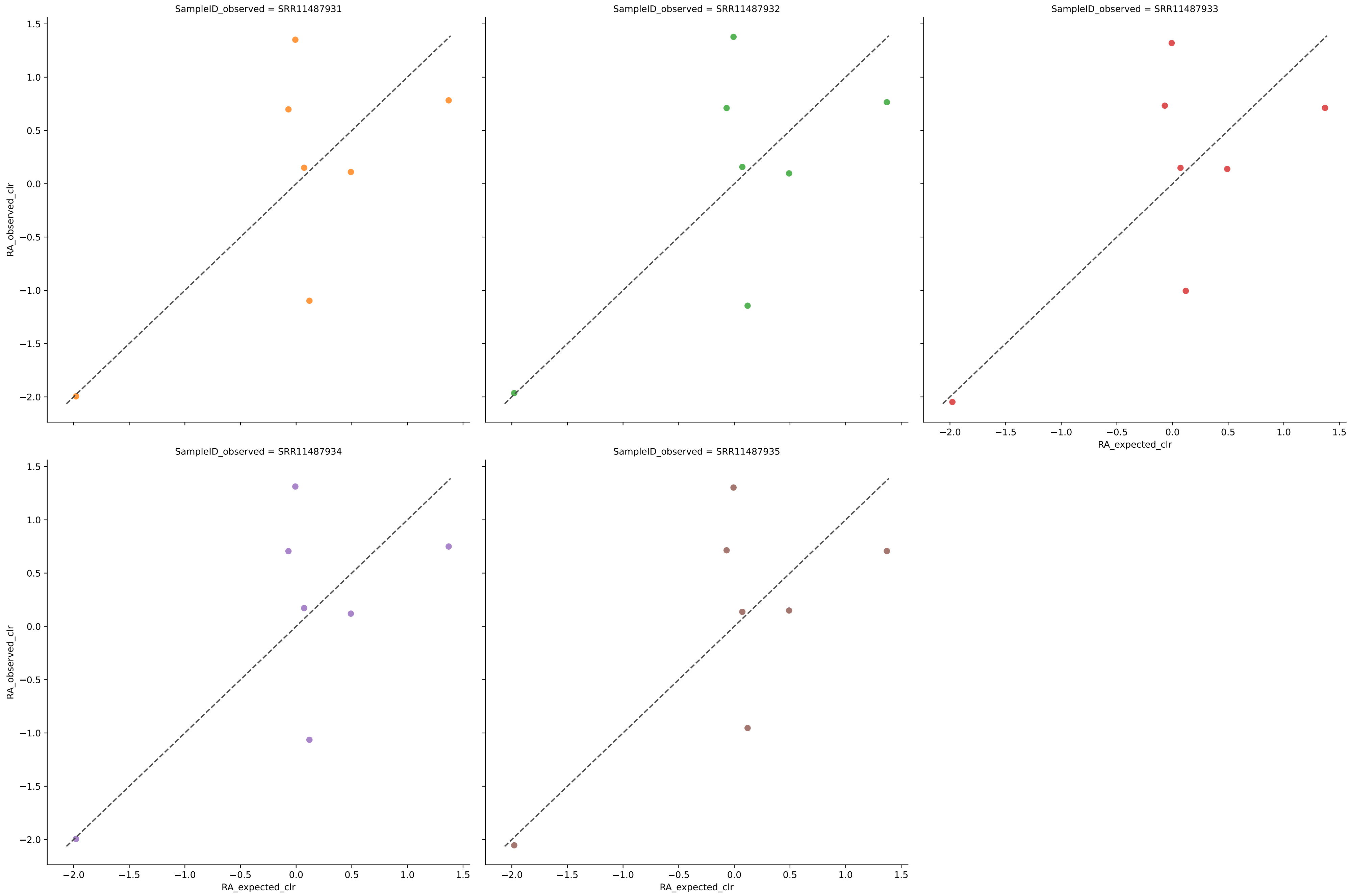
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487931	7	0.8412	0.0085	1.7999	0.8522	0.0103	100.0000	0.0000
SRR11487932	7	0.8301	0.0087	1.8762	0.8494	0.0106	100.0000	0.0000
SRR11487933	6	0.8130	0.0088	1.5454	0.8663	0.0104	100.0000	0.0000
SRR11487934	7	0.8346	0.0083	1.7049	0.8557	0.0102	100.0000	0.0000
SRR11487935	6	0.8257	0.0086	1.4585	0.8695	0.0101	100.0000	0.0000
Average	7	0.8289	0.0086	1.6770	0.8587	0.0103	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using wgsa in Experiment Amos hilo with filter 0.01



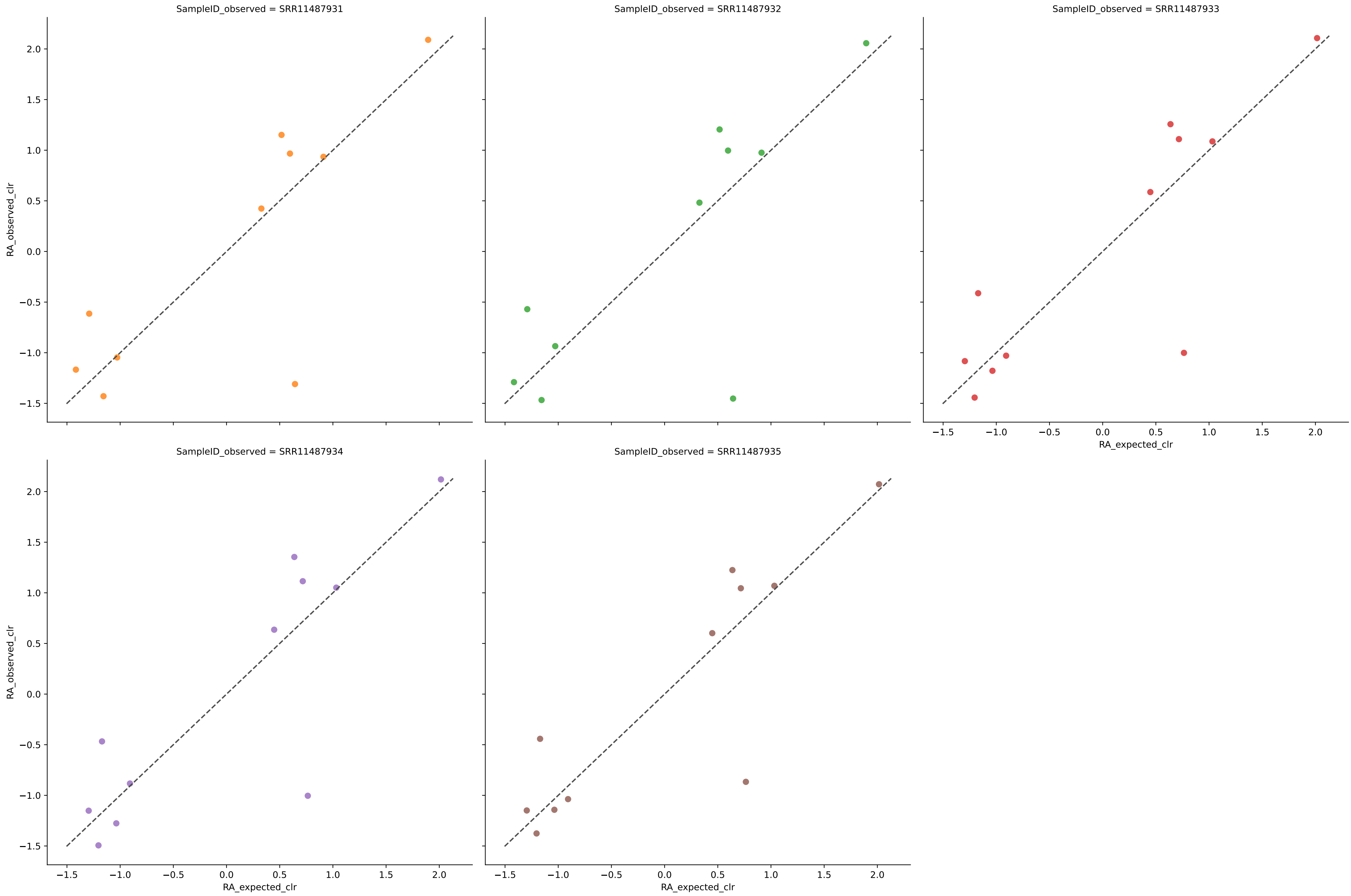
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487931	8	0.8976	0.0088	1.4706	0.8240	0.0114	100.0000	0.0000
SRR11487932	8	0.8939	0.0088	1.5407	0.8228	0.0114	100.0000	0.0000
SRR11487933	8	0.9009	0.0084	1.3341	0.8318	0.0105	100.0000	0.0000
SRR11487934	8	0.8976	0.0085	1.3870	0.8295	0.0108	100.0000	0.0000
SRR11487935	8	0.9057	0.0082	1.2581	0.8356	0.0103	100.0000	0.0000
Average	8	0.8991	0.0086	1.3981	0.8287	0.0109	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using wol in Experiment Amos hilo with filter 0.01



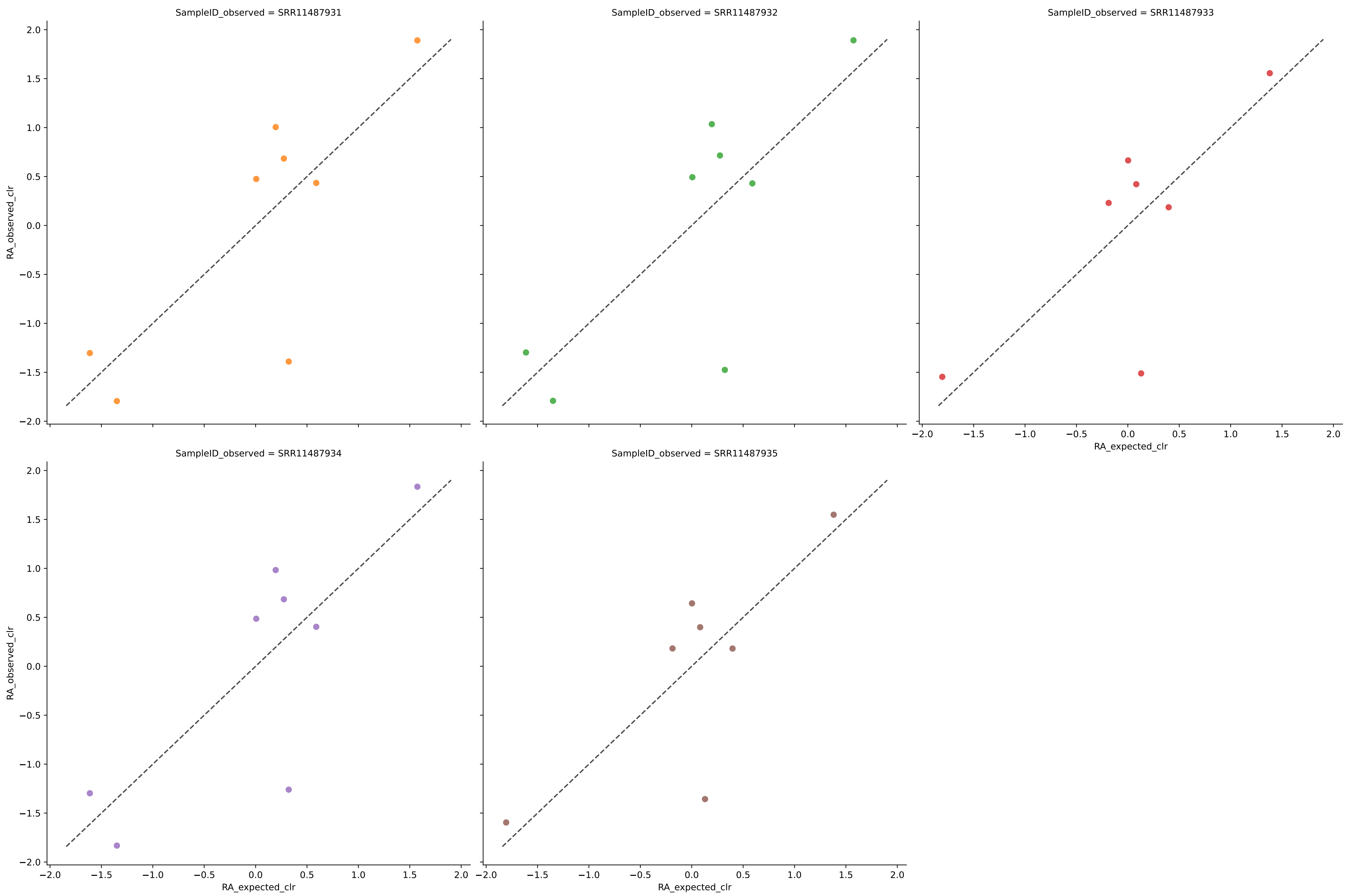
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487931	7	0.0799	0.0199	2.1011	0.6522	0.0266	100.0000	0.0000
SRR11487932	7	0.0670	0.0203	2.1578	0.6439	0.0273	100.0000	0.0000
SRR11487933	7	0.0644	0.0200	2.0592	0.6497	0.0269	100.0000	0.0000
SRR11487934	7	0.0782	0.0198	2.0681	0.6546	0.0264	100.0000	0.0000
SRR11487935	7	0.0684	0.0198	2.0124	0.6545	0.0266	100.0000	0.0000
Average	7	0.0716	0.0199	2.0797	0.6510	0.0268	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using bio4 in Experiment Amos hilo with filter 0.01



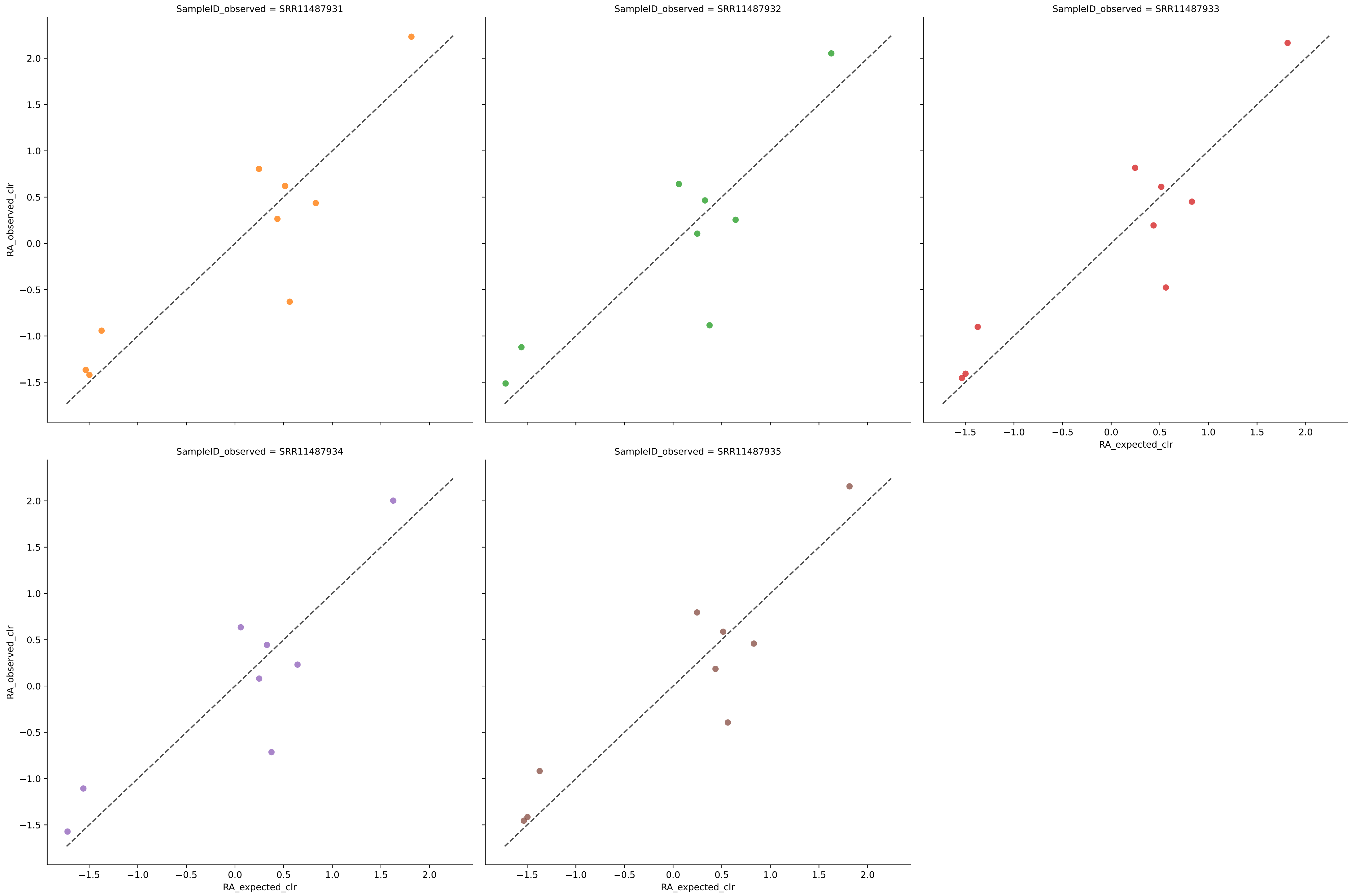
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487931	10	0.8908	0.0050	2.2353	0.8733	0.0078	100.0000	0.0000
SRR11487932	10	0.8747	0.0048	2.3922	0.8798	0.0080	100.0000	0.0000
SRR11487933	11	0.8851	0.0043	2.0969	0.8815	0.0072	100.0000	0.0000
SRR11487934	11	0.8702	0.0047	2.1221	0.8714	0.0077	100.0000	0.0000
SRR11487935	11	0.8907	0.0042	1.9366	0.8849	0.0070	100.0000	0.0000
Average	11	0.8823	0.0046	2.1566	0.8782	0.0075	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using jams in Experiment Amos hilo with filter 0.01



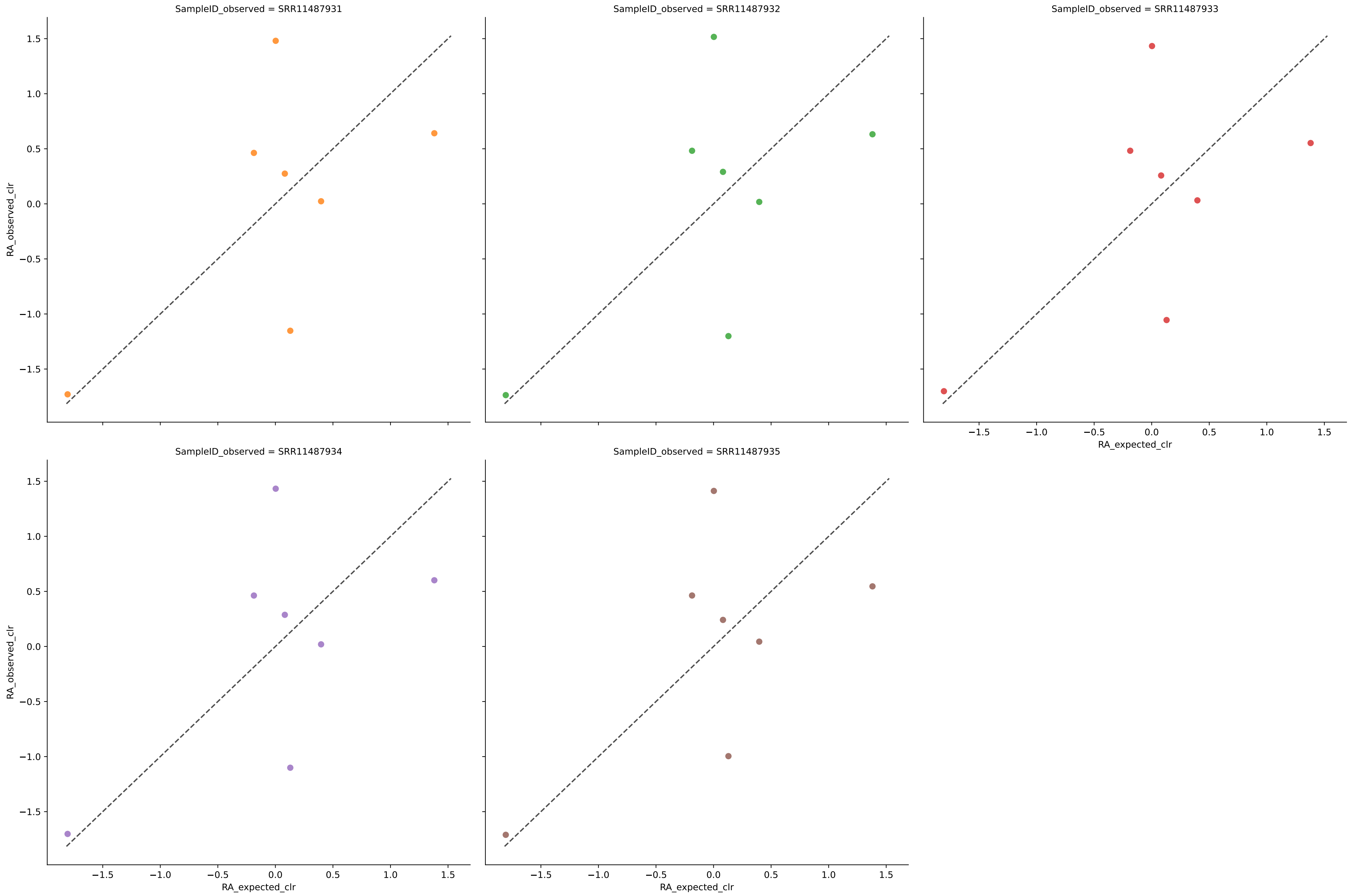
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487931	8	0.8543	0.0079	2.0959	0.8439	0.0101	100.0000	0.0000
SRR11487932	8	0.8445	0.0080	2.1884	0.8406	0.0104	100.0000	0.0000
SRR11487933	7	0.8393	0.0082	1.8871	0.8552	0.0102	100.0000	0.0000
SRR11487934	8	0.8472	0.0077	1.9892	0.8461	0.0100	100.0000	0.0000
SRR11487935	7	0.8509	0.0080	1.7245	0.8595	0.0099	100.0000	0.0000
Average	8	0.8472	0.0080	1.9770	0.8490	0.0101	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using wgsa in Experiment Amos hilo with filter 0.01



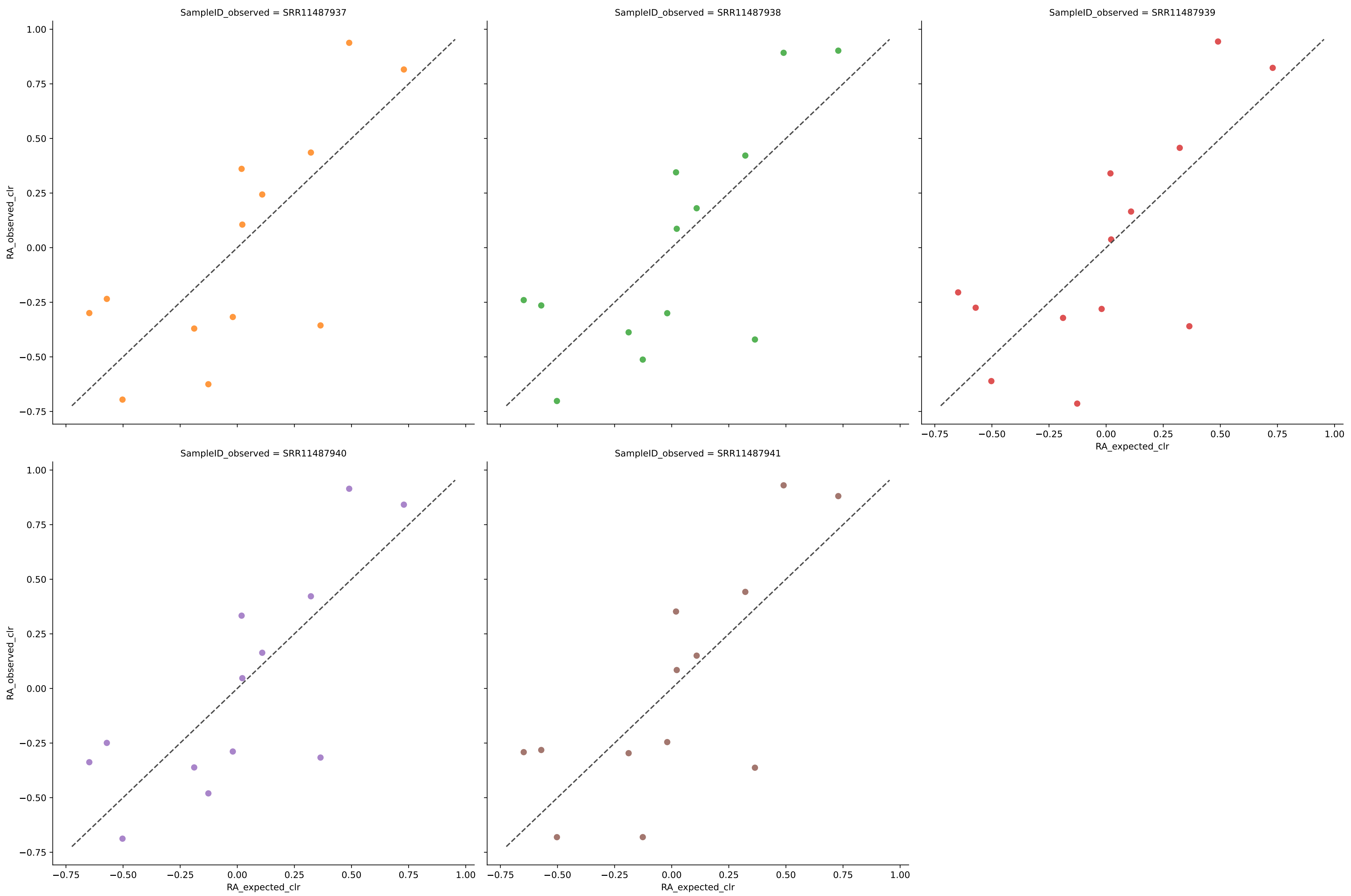
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487931	9	0.9024	0.0079	1.5246	0.8226	0.0116	100.0000	0.0000
SRR11487932	8	0.8989	0.0088	1.5913	0.8216	0.0122	100.0000	0.0000
SRR11487933	9	0.9063	0.0075	1.4054	0.8333	0.0106	100.0000	0.0000
SRR11487934	8	0.9017	0.0085	1.4486	0.8289	0.0116	100.0000	0.0000
SRR11487935	9	0.9108	0.0073	1.3261	0.8360	0.0104	100.0000	0.0000
Average	9	0.9040	0.0080	1.4592	0.8285	0.0113	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using wol in Experiment Amos hilo with filter 0.01



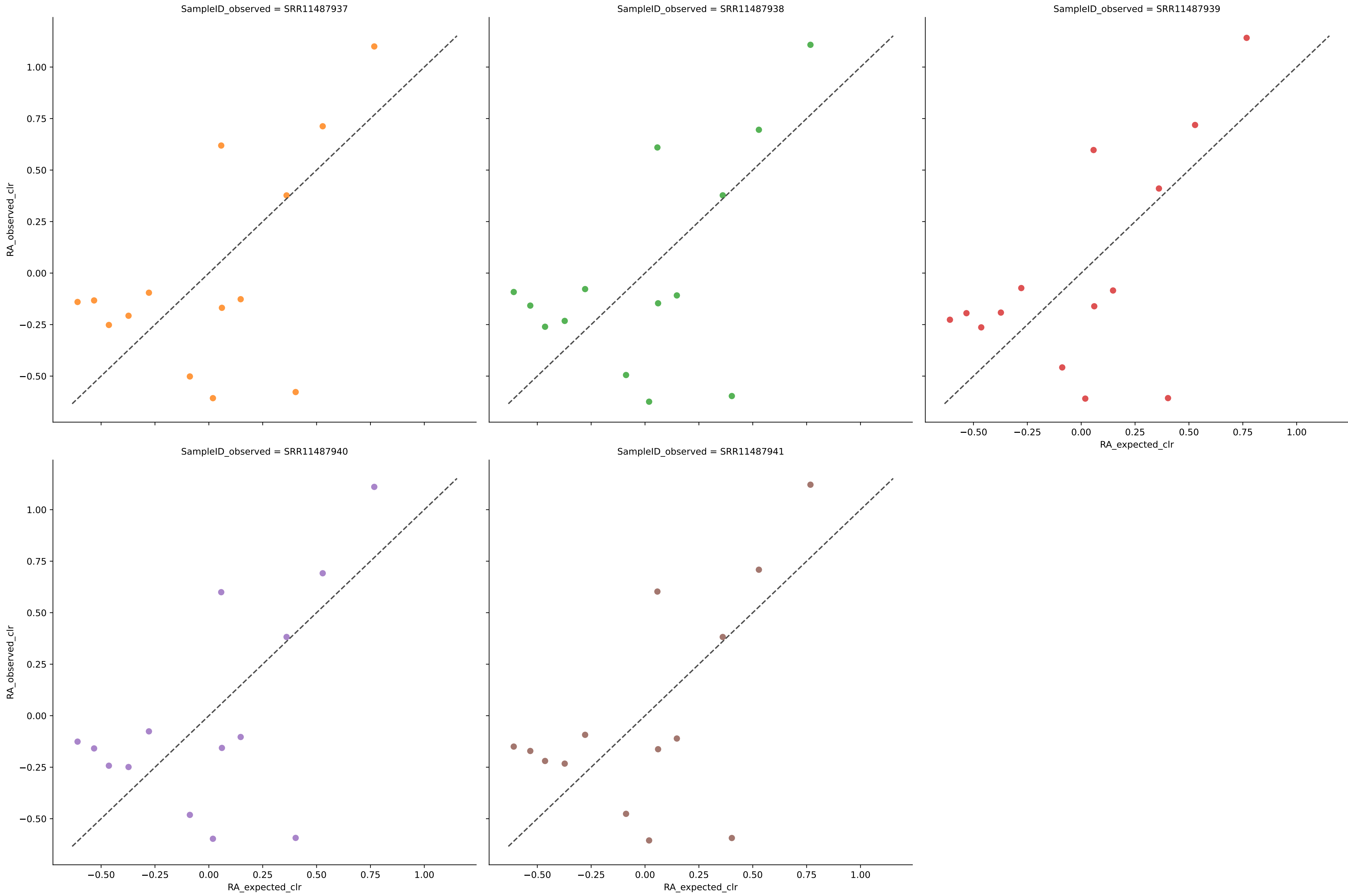
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487931	7	0.0304	0.0217	2.2311	0.6200	0.0306	100.0000	0.0000
SRR11487932	7	0.0243	0.0222	2.2935	0.6113	0.0313	100.0000	0.0000
SRR11487933	7	0.0216	0.0218	2.1813	0.6178	0.0306	100.0000	0.0000
SRR11487934	7	0.0292	0.0217	2.1872	0.6216	0.0302	100.0000	0.0000
SRR11487935	7	0.0236	0.0215	2.1289	0.6234	0.0303	100.0000	0.0000
Average	7	0.0258	0.0218	2.2044	0.6188	0.0306	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using bio4 in Experiment Amos mixed with filter 0.01



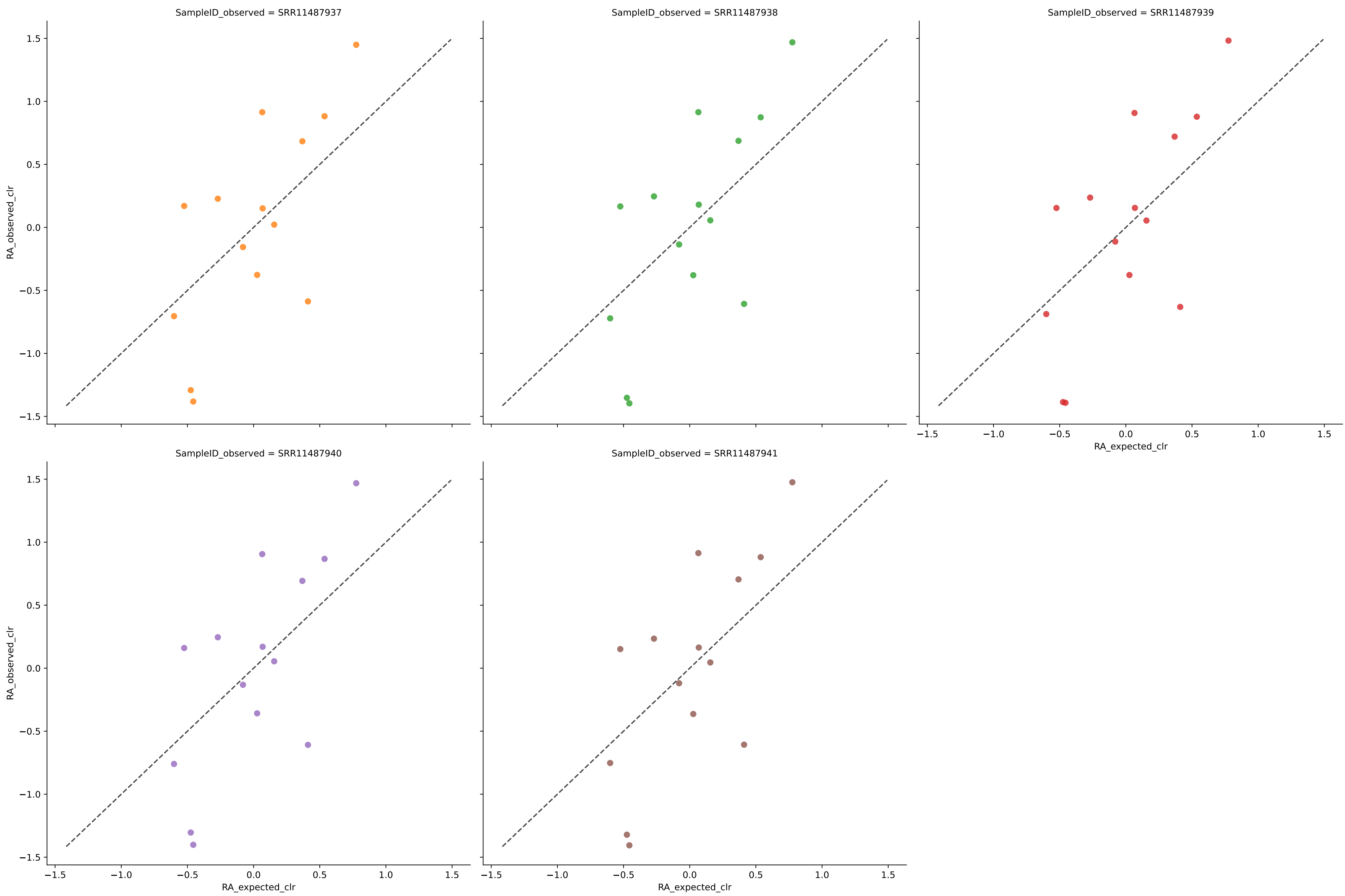
	Diversity	R ²	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487937	13	0.6380	0.0038	1.2346	0.8754	0.0051	100.0000	0.0000
SRR11487938	13	0.6683	0.0038	1.2249	0.8771	0.0050	100.0000	0.0000
SRR11487939	13	0.6319	0.0038	1.2611	0.8769	0.0052	100.0000	0.0000
SRR11487940	13	0.6775	0.0035	1.1091	0.8853	0.0048	100.0000	0.0000
SRR11487941	13	0.6683	0.0037	1.2201	0.8789	0.0051	100.0000	0.0000
Average	13	0.6568	0.0037	1.2100	0.8787	0.0050	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using jams in Experiment Amos mixed with filter 0.01



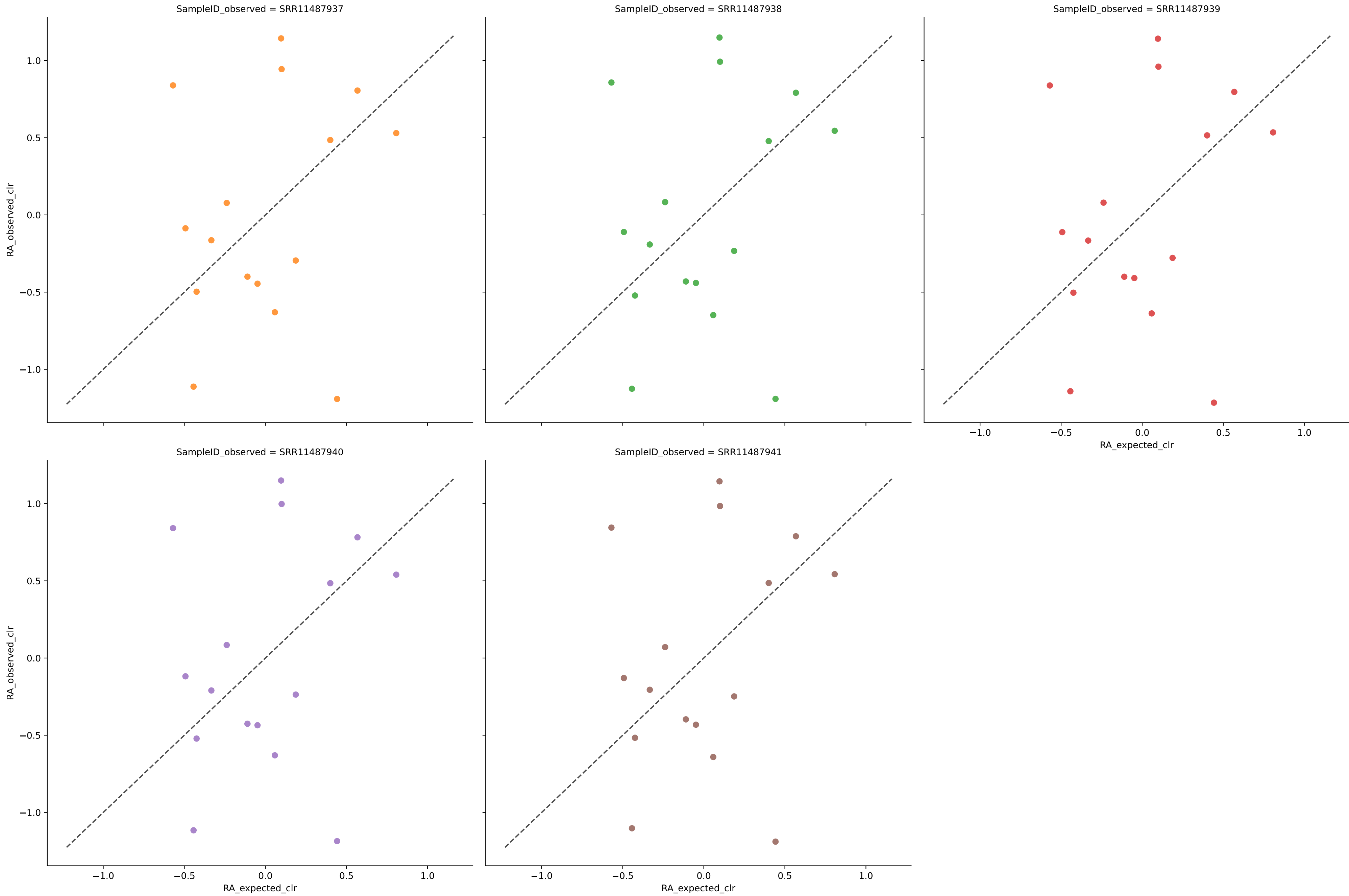
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487937	14	0.5414	0.0046	1.6117	0.8392	0.0057	100.0000	0.0000
SRR11487938	14	0.5390	0.0046	1.6265	0.8397	0.0058	100.0000	0.0000
SRR11487939	14	0.5698	0.0044	1.5811	0.8444	0.0057	100.0000	0.0000
SRR11487940	14	0.5476	0.0045	1.5980	0.8418	0.0057	100.0000	0.0000
SRR11487941	14	0.5522	0.0046	1.6001	0.8407	0.0058	100.0000	0.0000
Average	14	0.5500	0.0045	1.6035	0.8412	0.0058	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using wgsa in Experiment Amos mixed with filter 0.01



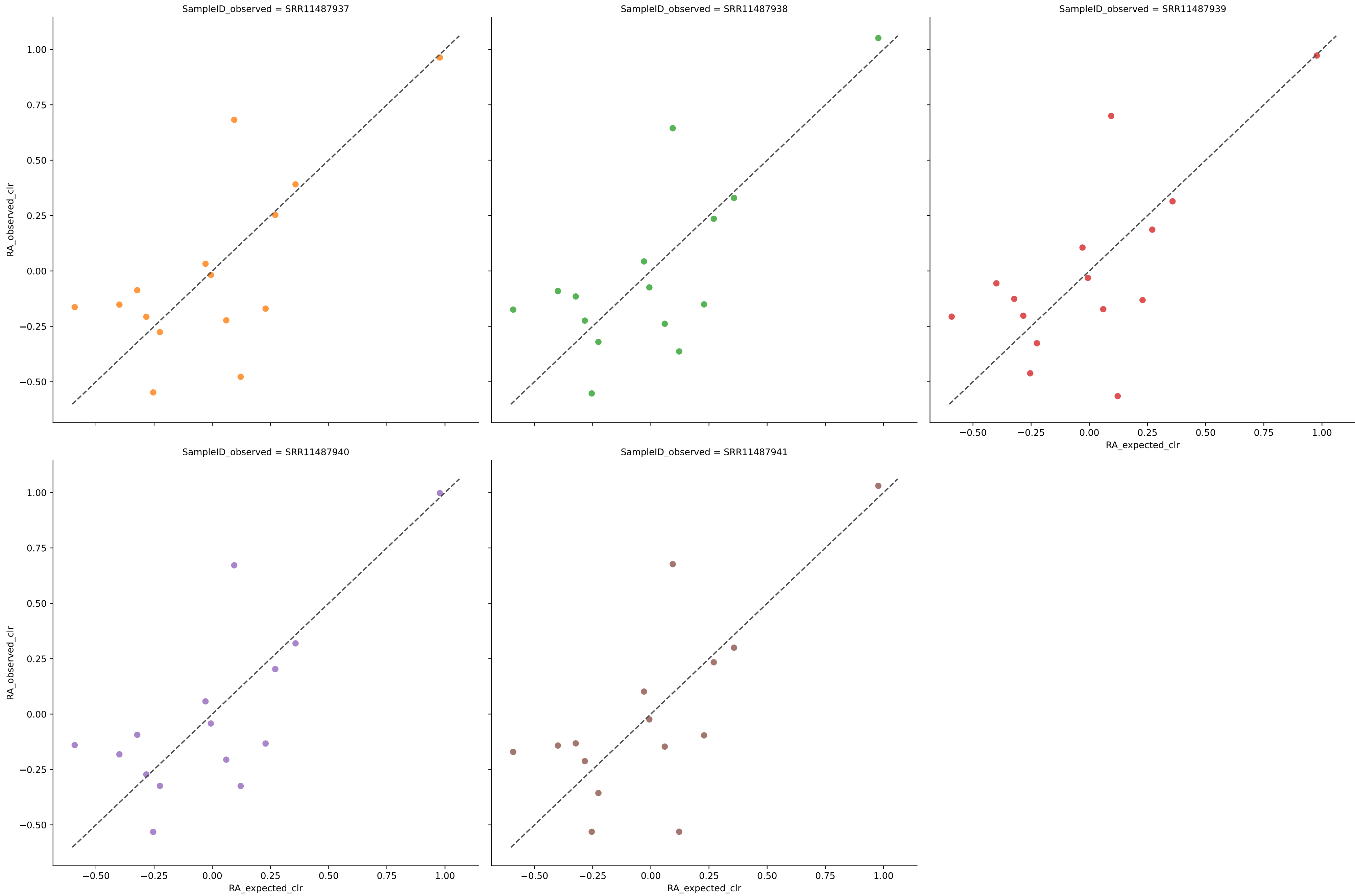
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487937	14	0.6079	0.0060	2.2021	0.7890	0.0076	100.0000	0.0000
SRR11487938	14	0.6085	0.0060	2.2487	0.7891	0.0077	100.0000	0.0000
SRR11487939	14	0.6129	0.0061	2.2696	0.7872	0.0077	100.0000	0.0000
SRR11487940	14	0.6117	0.0060	2.2255	0.7896	0.0076	100.0000	0.0000
SRR11487941	14	0.6147	0.0061	2.2373	0.7880	0.0077	100.0000	0.0000
Average	14	0.6112	0.0060	2.2366	0.7886	0.0077	100.0000	0.0000

Expected vs. Observed Relative Abundance for genus using wol in Experiment Amos mixed with filter 0.01



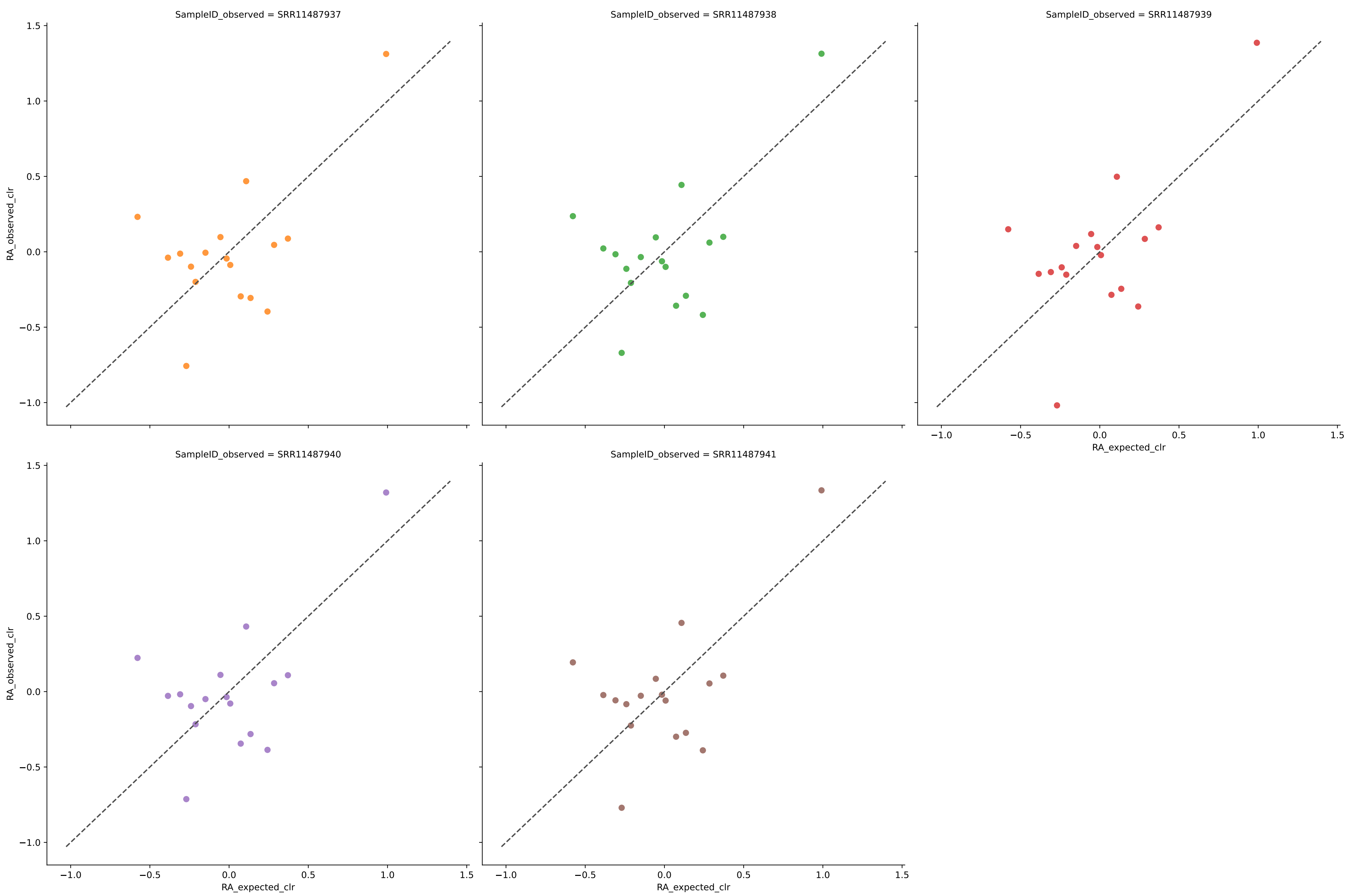
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487937	16	0.0645	0.0066	2.8809	0.7364	0.0087	100.0000	0.0000
SRR11487938	16	0.0633	0.0066	2.9006	0.7340	0.0089	100.0000	0.0000
SRR11487939	16	0.0662	0.0065	2.8965	0.7387	0.0087	100.0000	0.0000
SRR11487940	16	0.0643	0.0066	2.8819	0.7351	0.0088	100.0000	0.0000
SRR11487941	16	0.0658	0.0066	2.8748	0.7369	0.0088	100.0000	0.0000
Average	16	0.0648	0.0066	2.8869	0.7362	0.0088	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using bio4 in Experiment Amos mixed with filter 0.01



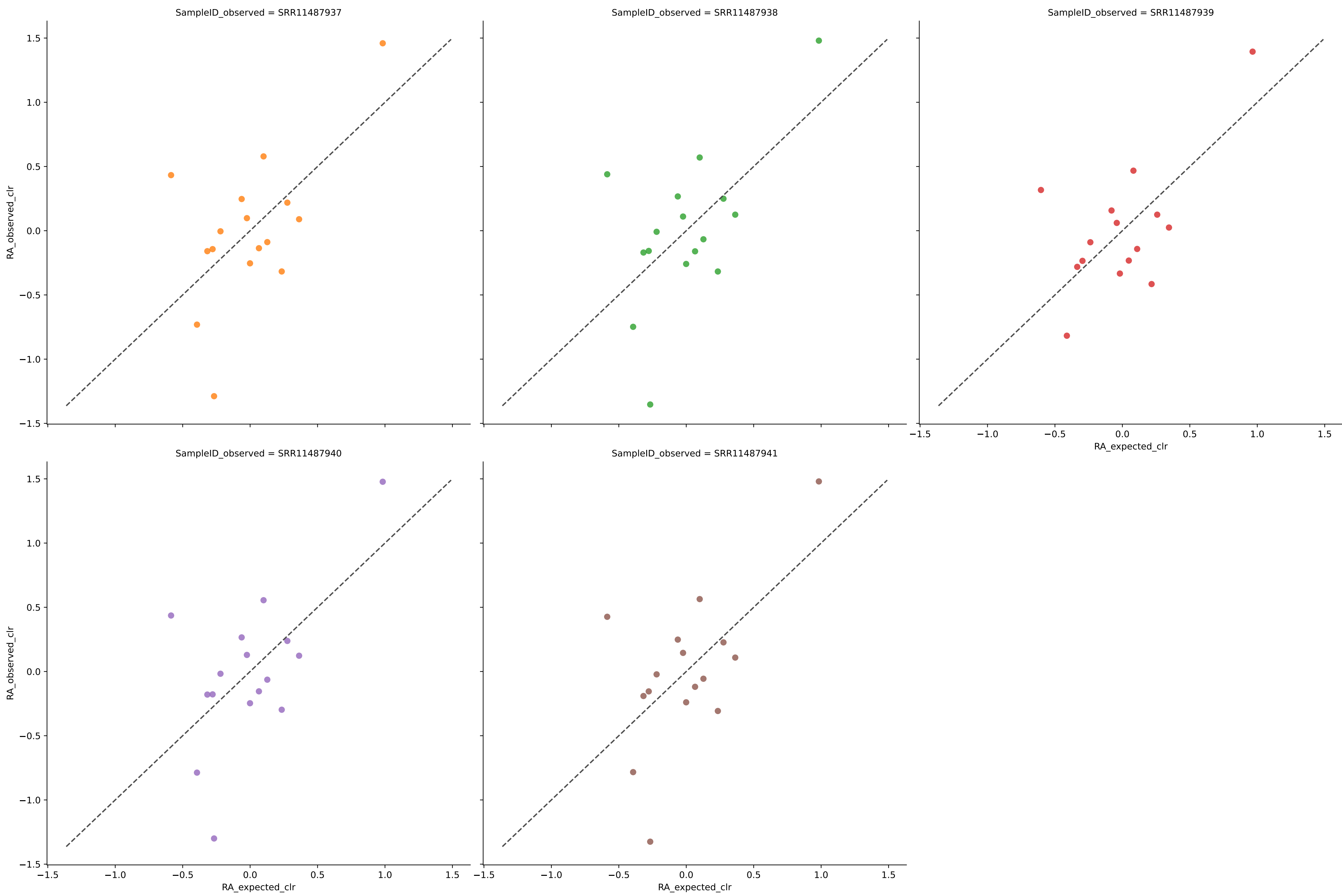
	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487937	15	0.6641	0.0027	1.1599	0.9002	0.0038	100.0000	0.0000
SRR11487938	15	0.7255	0.0027	1.0966	0.8976	0.0036	100.0000	0.0000
SRR11487939	15	0.6459	0.0028	1.1898	0.8938	0.0040	100.0000	0.0000
SRR11487940	15	0.7012	0.0026	1.0679	0.9033	0.0036	100.0000	0.0000
SRR11487941	15	0.6991	0.0027	1.1479	0.8974	0.0038	100.0000	0.0000
Average	15	0.6872	0.0027	1.1324	0.8985	0.0038	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using jams in Experiment Amos mixed with filter 0.01



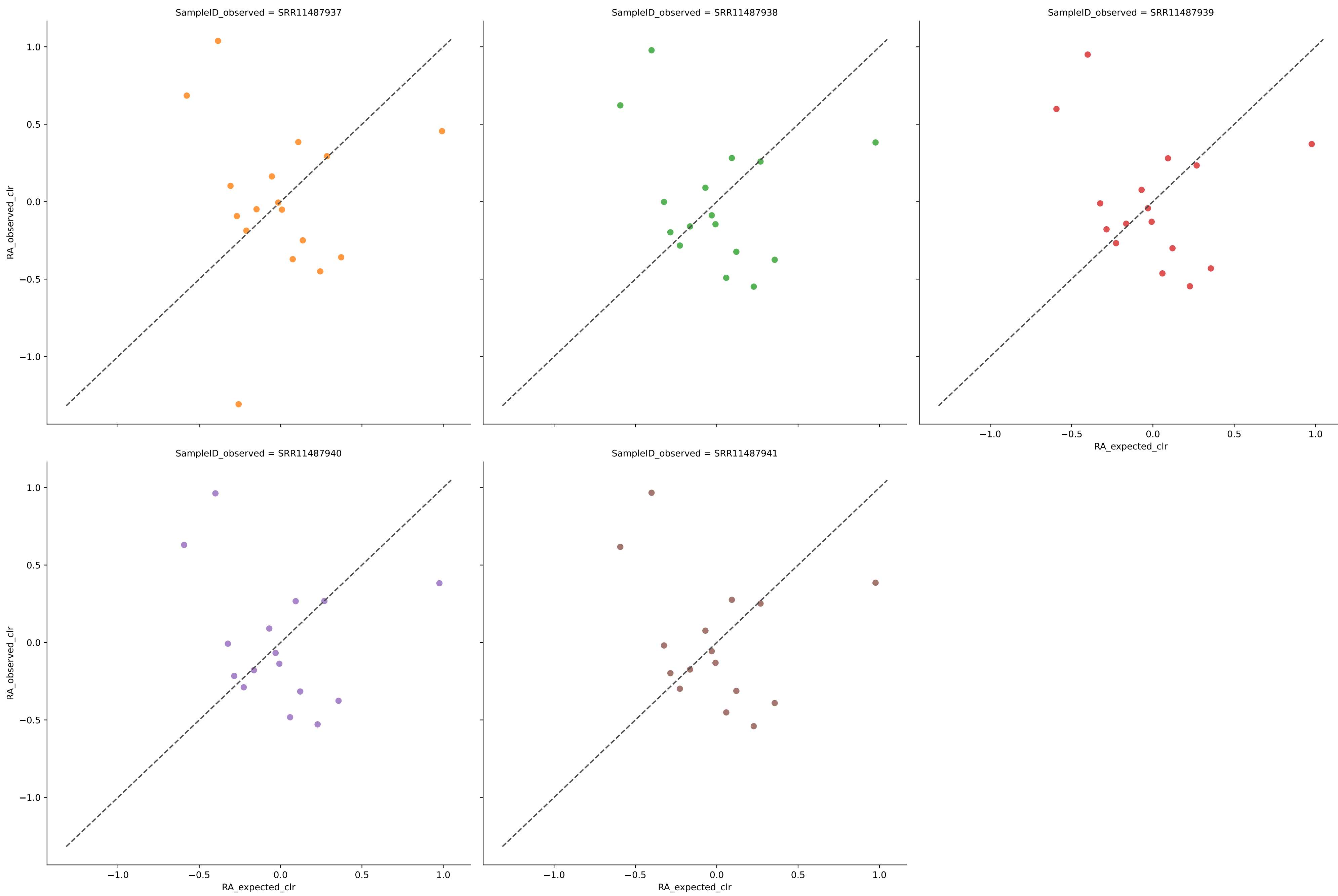
	Diversity	R ²	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487937	17	0.6678	0.0035	1.5112	0.8526	0.0043	100.0000	0.0000
SRR11487938	17	0.6652	0.0035	1.5122	0.8521	0.0044	100.0000	0.0000
SRR11487939	17	0.7225	0.0032	1.5163	0.8603	0.0043	100.0000	0.0000
SRR11487940	17	0.6796	0.0034	1.4818	0.8550	0.0043	100.0000	0.0000
SRR11487941	17	0.6900	0.0034	1.4737	0.8557	0.0043	100.0000	0.0000
Average	17	0.6850	0.0034	1.4990	0.8551	0.0043	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using wgsa in Experiment Amos mixed with filter 0.01



	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487937	16	0.6864	0.0042	1.8398	0.8312	0.0055	100.0000	0.0000
SRR11487938	16	0.6953	0.0043	1.8839	0.8310	0.0056	100.0000	0.0000
SRR11487939	15	0.6980	0.0041	1.4839	0.8421	0.0056	100.0000	0.0000
SRR11487940	16	0.6998	0.0042	1.8456	0.8317	0.0056	100.0000	0.0000
SRR11487941	16	0.7008	0.0042	1.8543	0.8321	0.0056	100.0000	0.0000
Average	16	0.6961	0.0042	1.7815	0.8336	0.0056	100.0000	0.0000

Expected vs. Observed Relative Abundance for species using wol in Experiment Amos mixed with filter 0.01



	Diversity	R ²	MAE	AD	1-BC	RMSE	Sens	FPRA
SRR11487937	17	0.0008	0.0056	2.5896	0.7664	0.0082	100.0000	0.0000
SRR11487938	16	0.0072	0.0056	2.3587	0.7754	0.0084	100.0000	0.0000
SRR11487939	16	0.0080	0.0055	2.3366	0.7800	0.0083	100.0000	0.0000
SRR11487940	16	0.0065	0.0055	2.3415	0.7782	0.0083	100.0000	0.0000
SRR11487941	16	0.0062	0.0055	2.3355	0.7787	0.0083	100.0000	0.0000
Average	16	0.0057	0.0055	2.3924	0.7757	0.0083	100.0000	0.0000