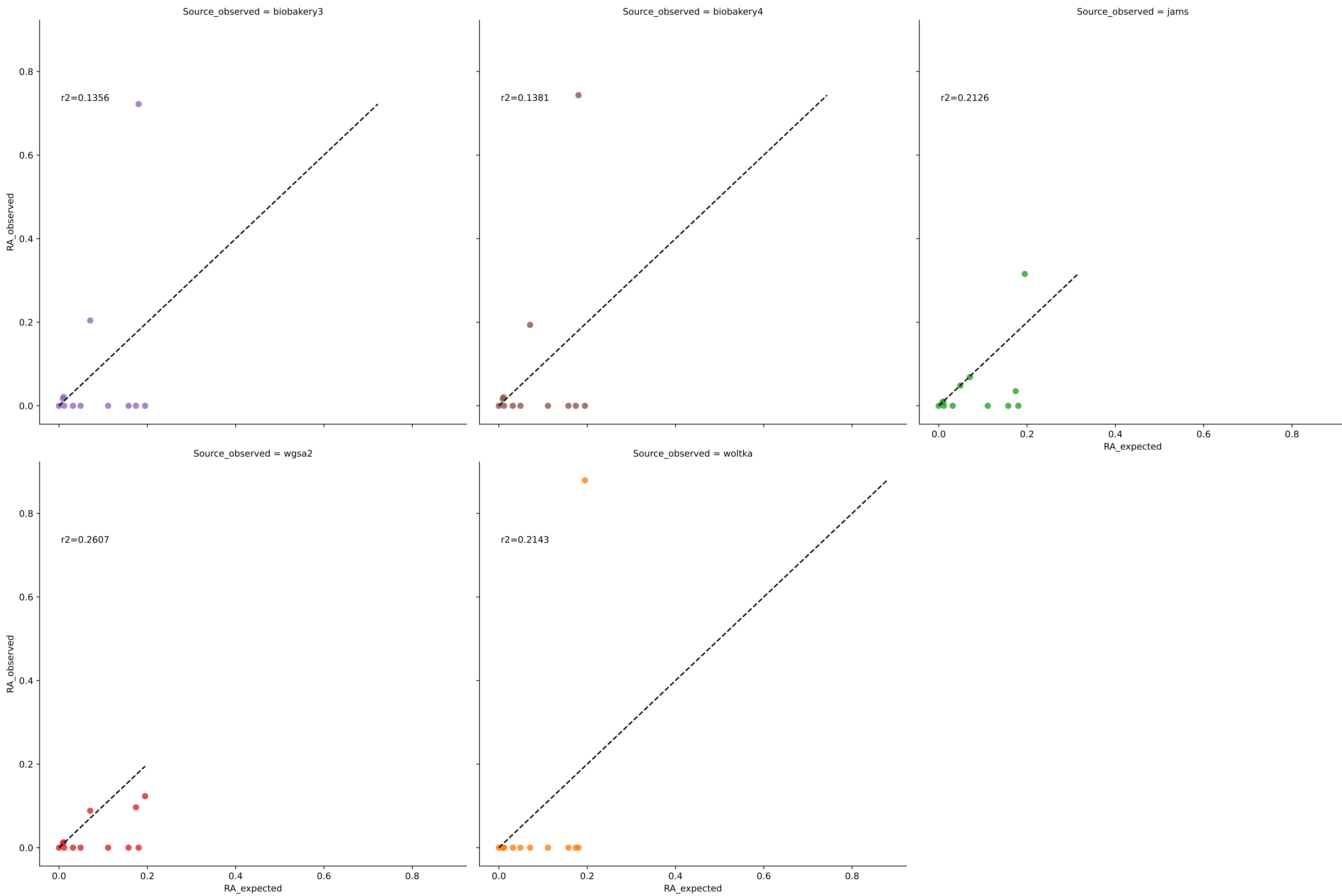
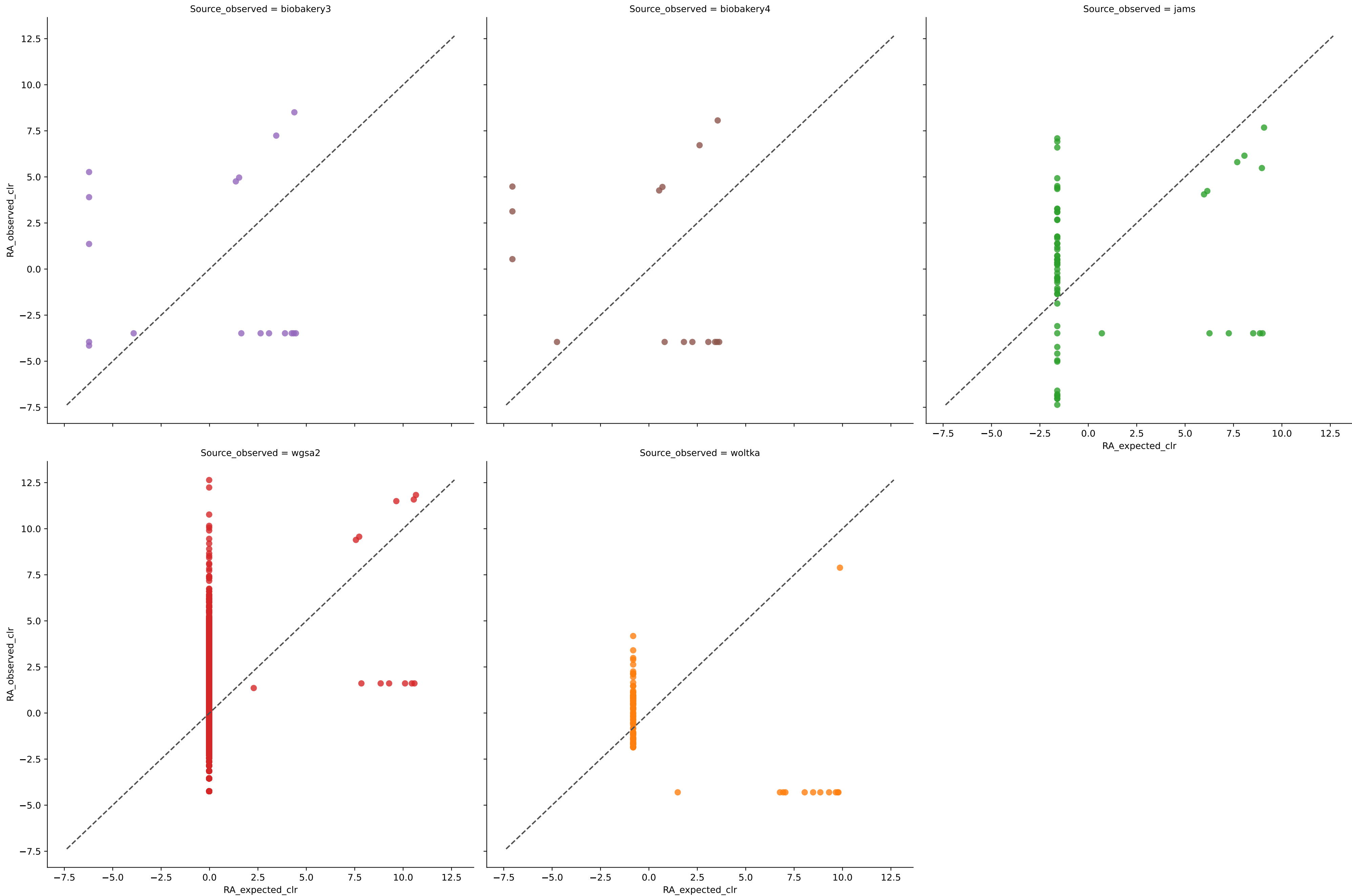


Bivariate Linear Regression for Sample S1 in Experiment bmock12 (Species at filter threshold 0)

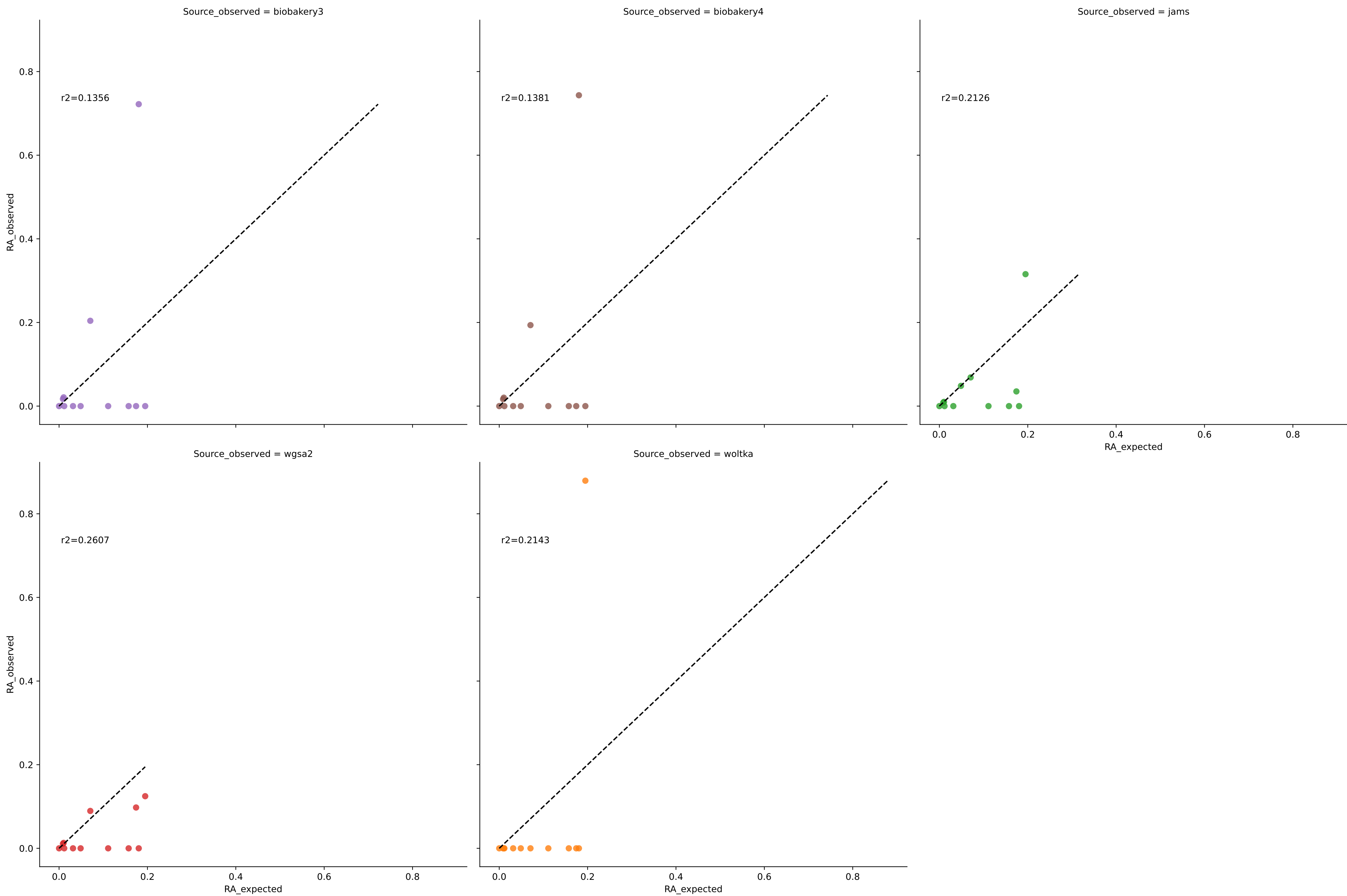


Bivariate Linear Regression for Sample S1 in Experiment bmock12 (Species at filter threshold 0)

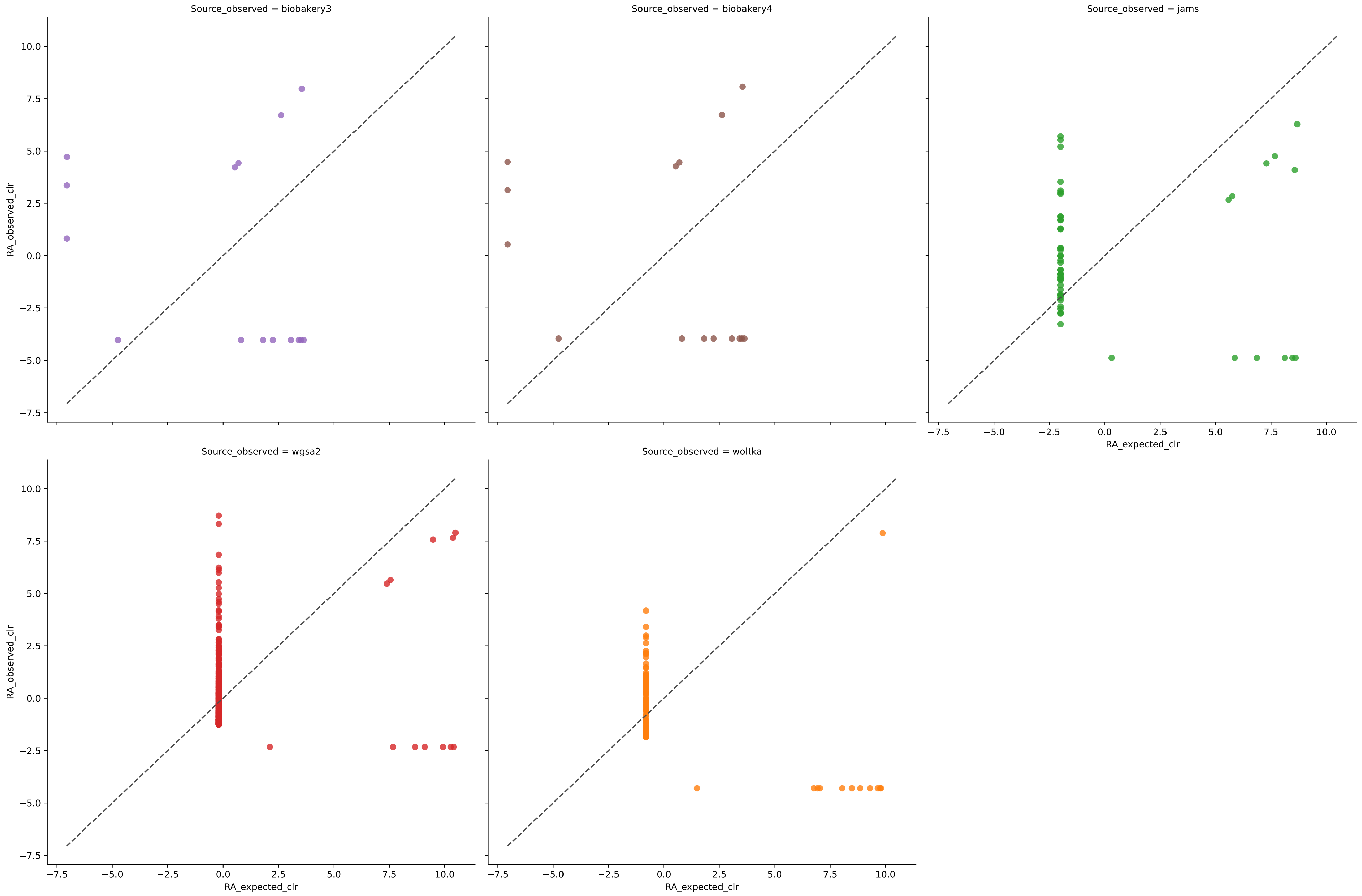


	Diversity	R ²	MAE	AD	1-BC	RMSE	Sens	FPRA
biobakery3	17	0.1670	0.0859	26.4923	0.2702	0.1574	33.3333	3.6049
biobakery4	15	0.1578	0.0973	25.9132	0.2702	0.1716	33.3333	2.6303
jams	66	0.1721	0.0192	39.5470	0.3655	0.0508	50.0000	51.3979
wgsa2	5631	0.1031	0.0002	177.6086	0.3097	0.0060	50.0000	66.8175
woltka	130	0.2516	0.0124	44.8238	0.1949	0.0667	8.3333	12.0865

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

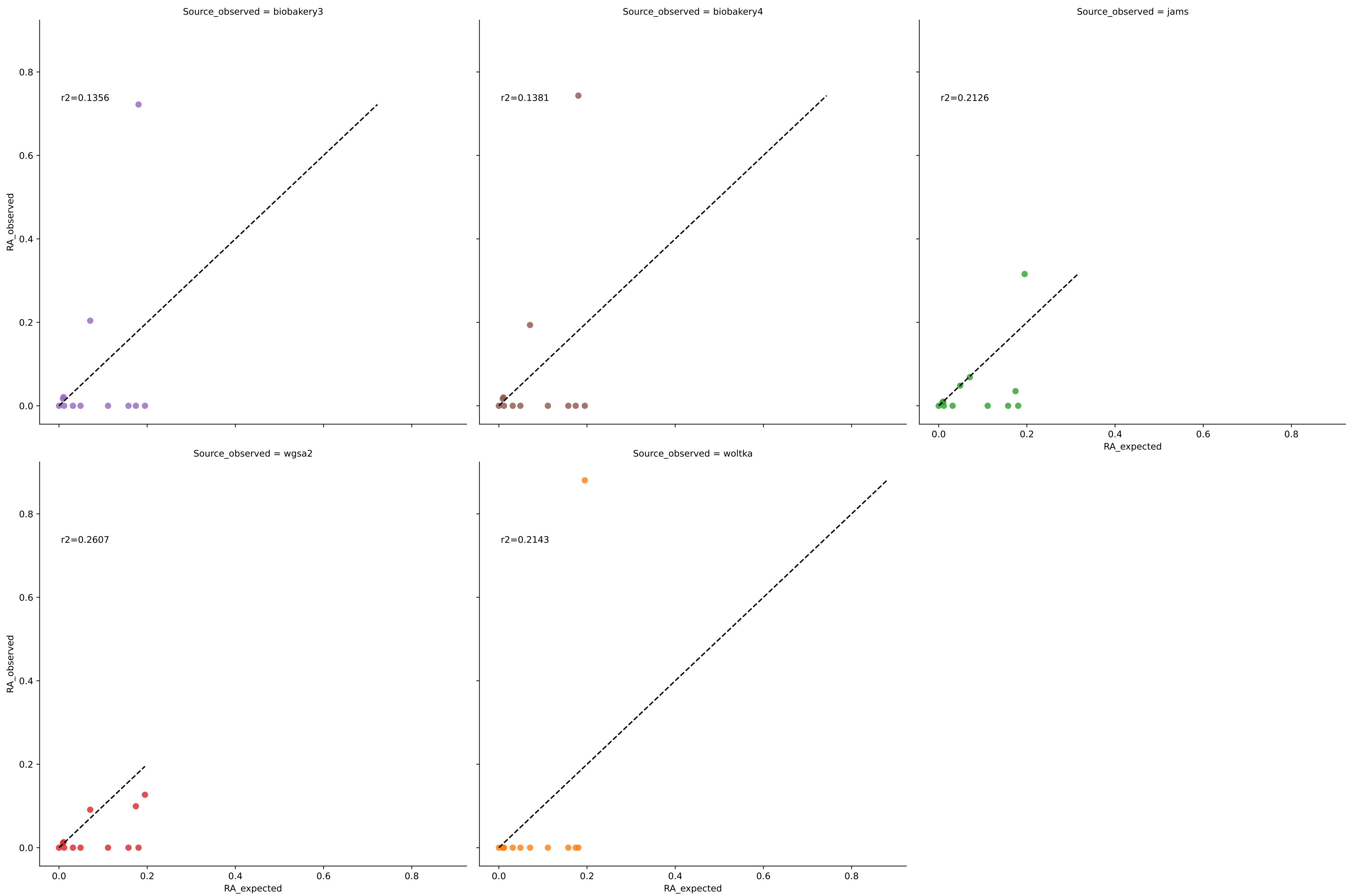


Bivariate Linear Regression for Sample S1 in Experiment bmock12 (Species at filter threshold 1e-05)

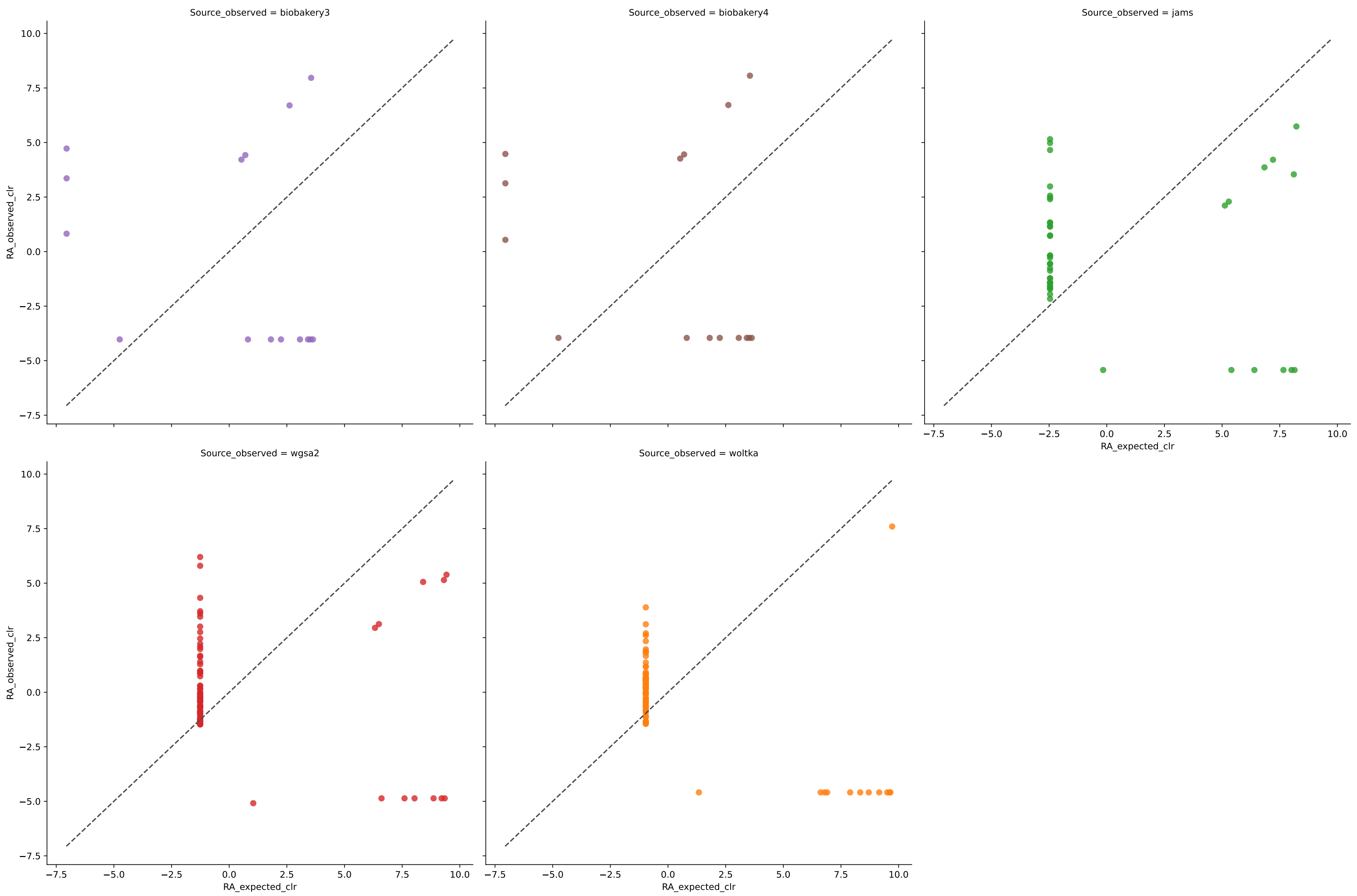


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
biobakery3	15	0.1539	0.0973	26.2902	0.2702	0.1676	33.3333	3.6044
biobakery4	15	0.1578	0.0973	25.9132	0.2702	0.1716	33.3333	2.6303
jams	53	0.1596	0.0239	35.5864	0.3655	0.0566	50.0000	51.3971
wgsa2	551	0.0983	0.0025	43.6657	0.3121	0.0191	41.6667	66.4502
woltka	130	0.2516	0.0124	44.8238	0.1949	0.0667	8.3333	12.0865

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

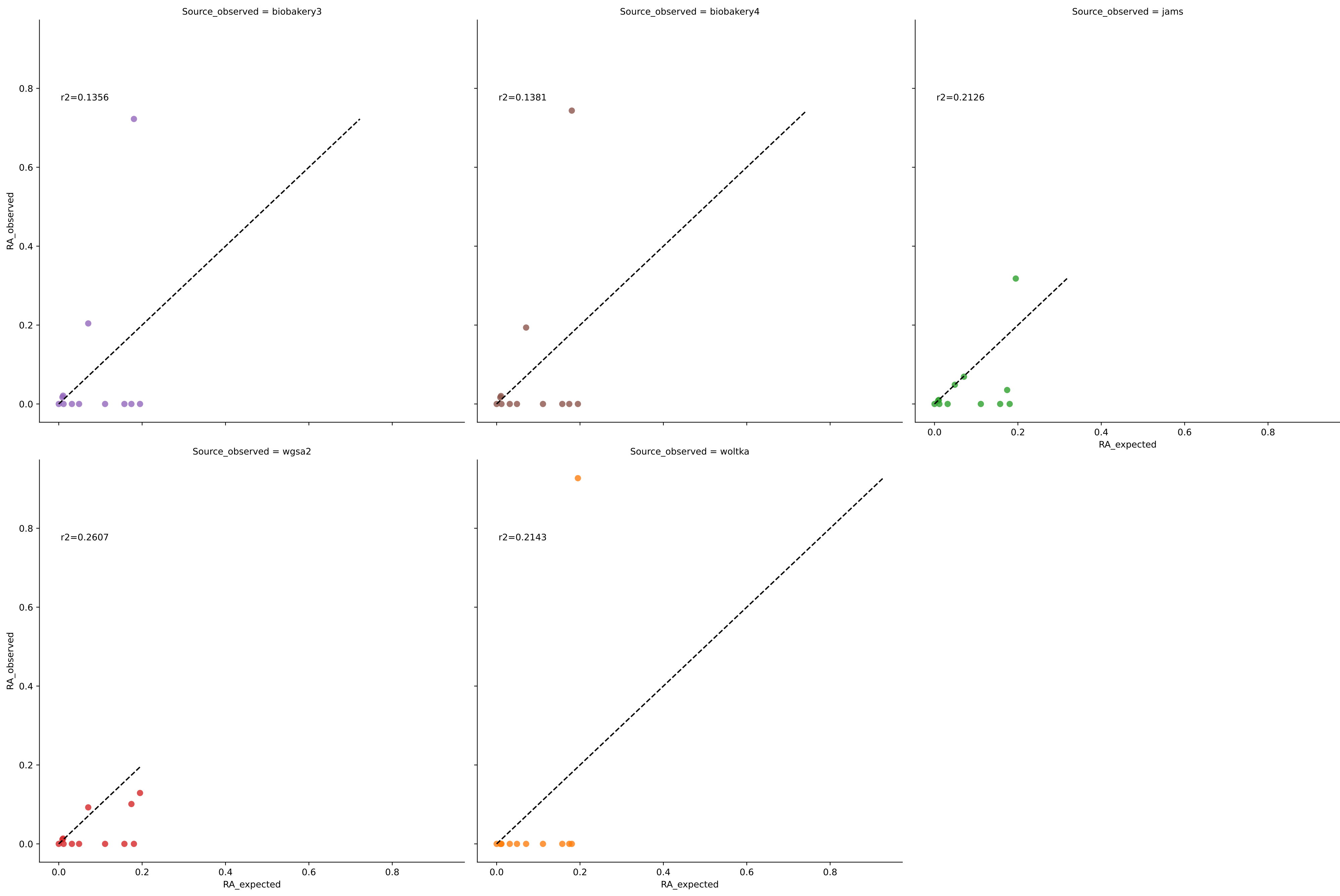


Bivariate Linear Regression for Sample S1 in Experiment bmock12 (Species at filter threshold 0.0001)

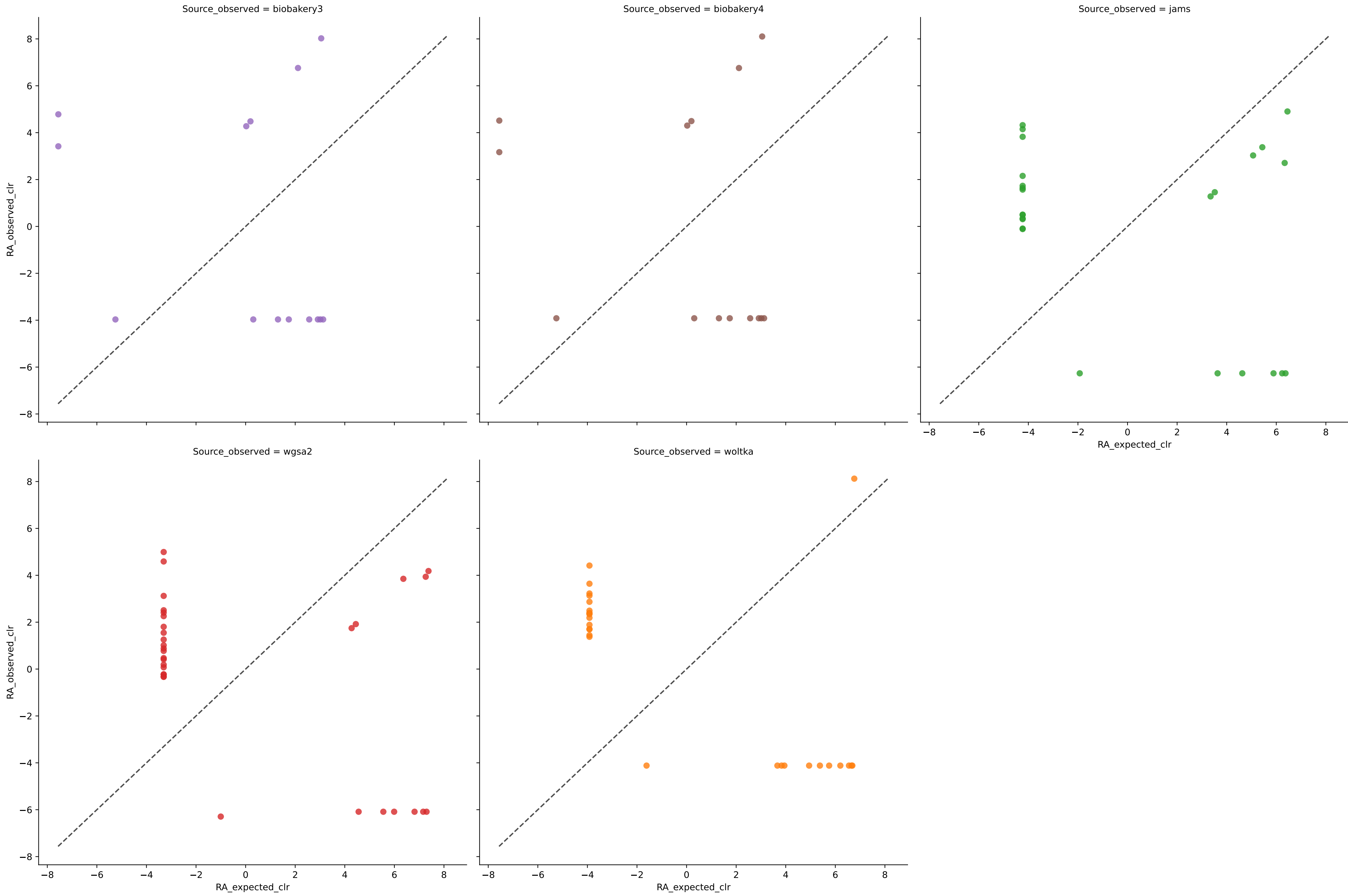


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
biobakery3	15	0.1539	0.0973	26.2902	0.2702	0.1676	33.3333	3.6044
biobakery4	15	0.1578	0.0973	25.9132	0.2702	0.1716	33.3333	2.6303
jams	43	0.1446	0.0295	35.5374	0.3656	0.0629	50.0000	51.3670
wgsa2	84	0.0697	0.0163	38.9395	0.3159	0.0494	50.0000	65.8720
woltka	110	0.2503	0.0146	44.6628	0.1949	0.0726	8.3333	11.9647

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

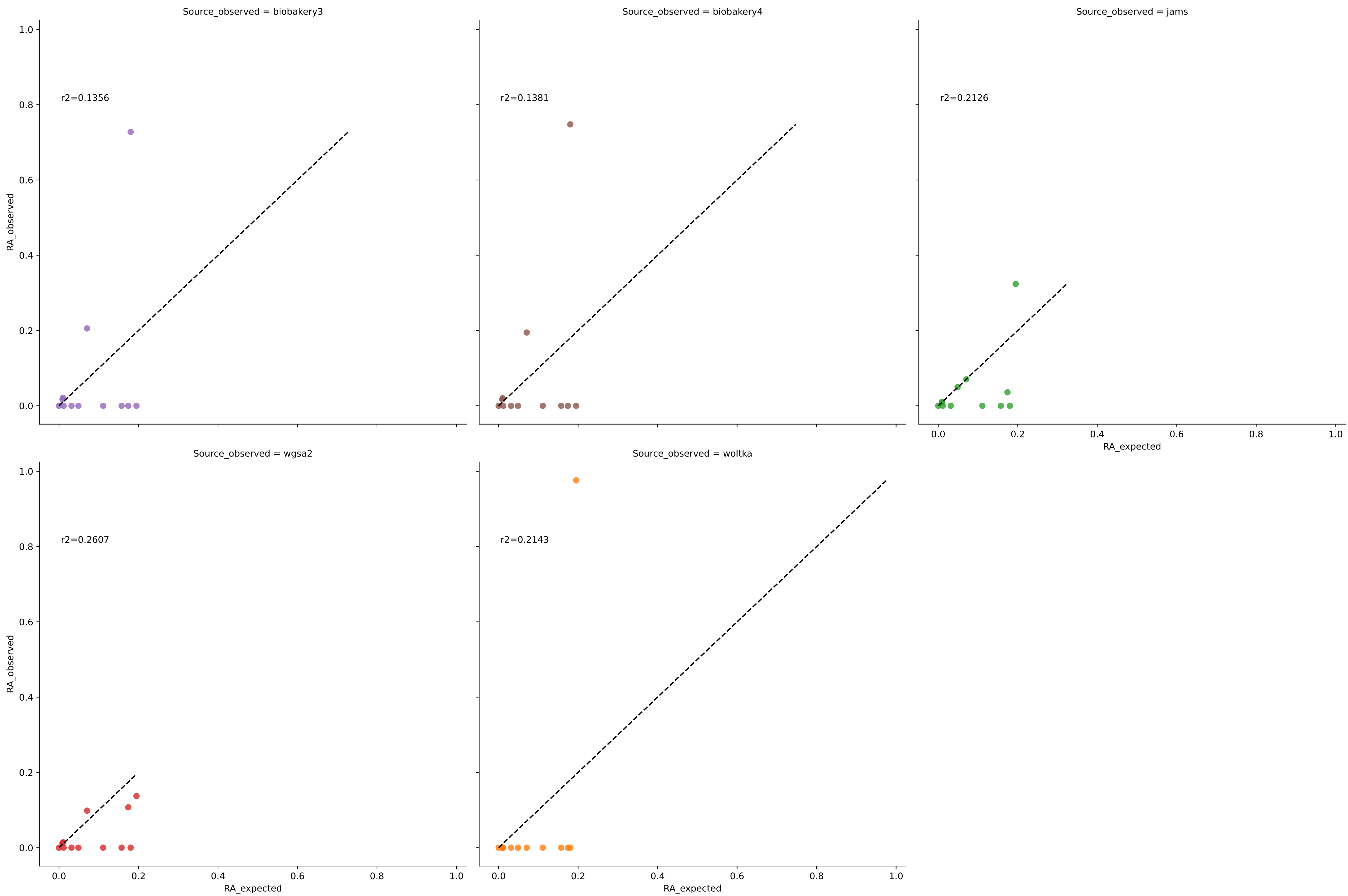


Bivariate Linear Regression for Sample S1 in Experiment bmock12 (Species at filter threshold 0.001)

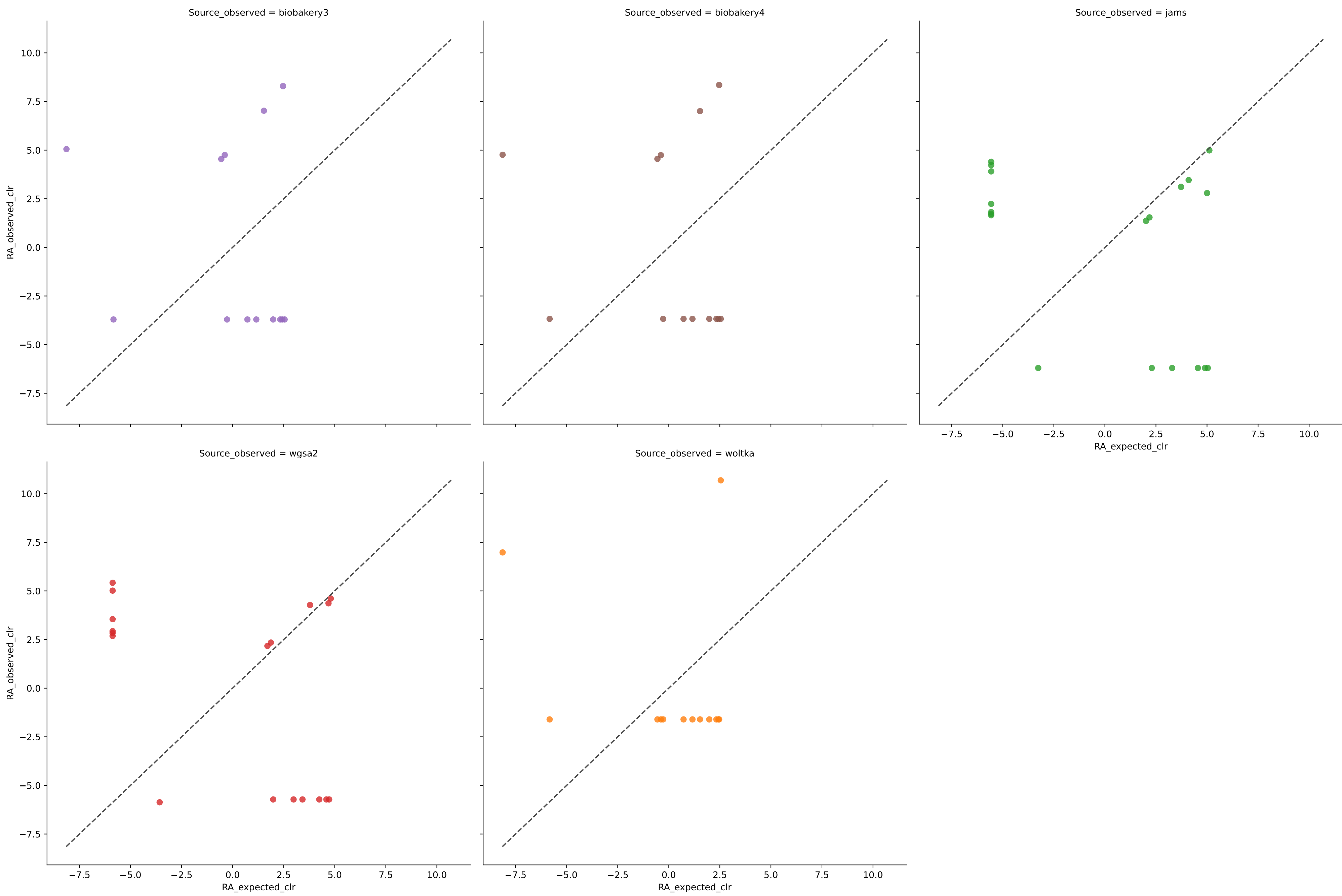


	Diversity	R ²	MAE	AD	1-BC	RMSE	Sens	FPRA
biobakery3	14	0.1458	0.1043	24.9967	0.2702	0.1736	33.3333	3.5494
biobakery4	14	0.1502	0.1043	24.6946	0.2702	0.1777	33.3333	2.5913
jams	25	0.0872	0.0507	34.7101	0.3668	0.0828	50.0000	51.0282
wgsa2	32	0.0222	0.0425	38.2831	0.3199	0.0808	50.0000	65.2799
woltka	27	0.2286	0.0596	39.2860	0.1949	0.1546	8.3333	7.3008

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

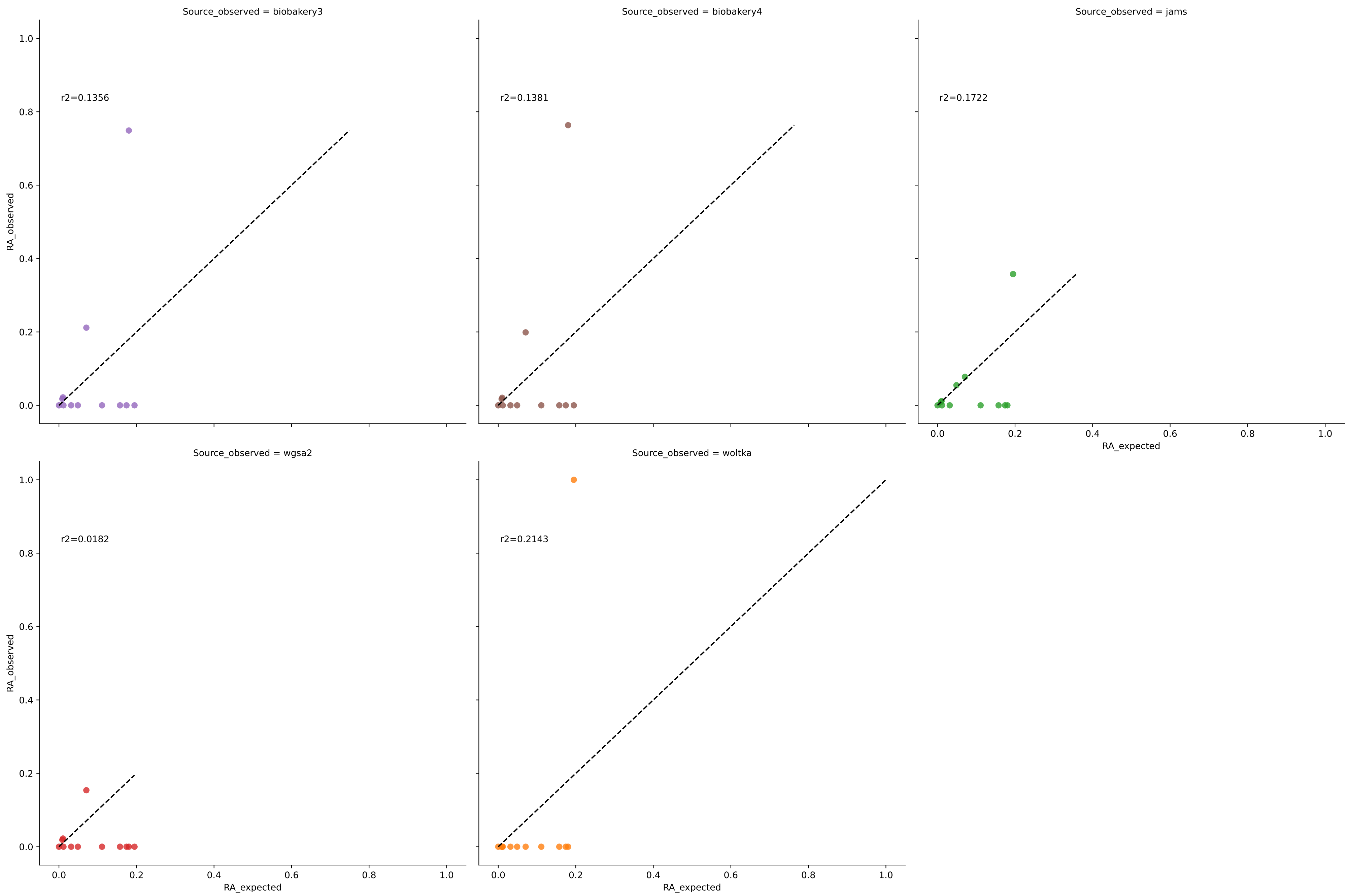


Bivariate Linear Regression for Sample S1 in Experiment bmock12 (Species at filter threshold 0.01)

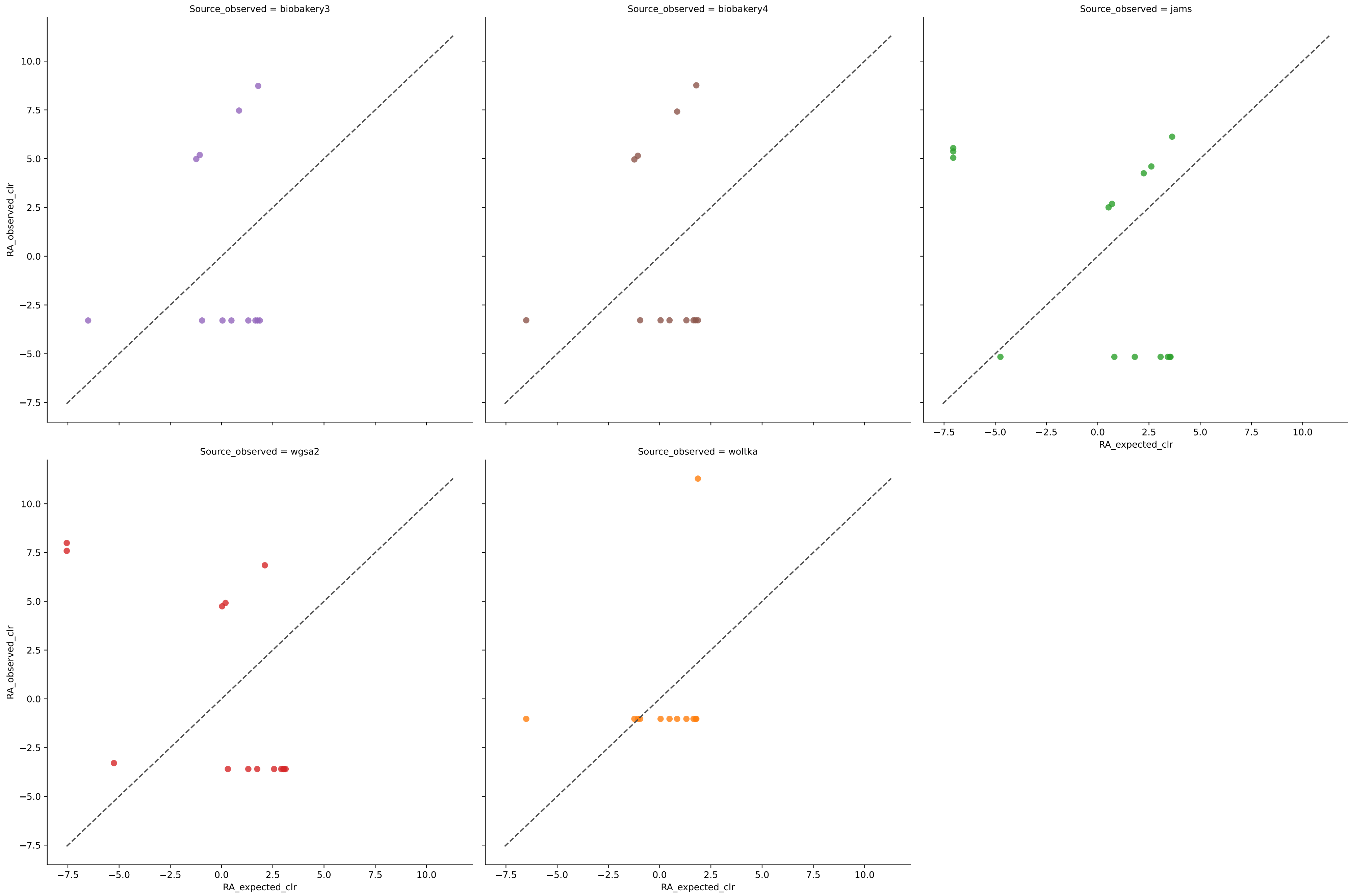


	Diversity	R ²	MAE	AD	1-BC	RMSE	Sens	FPRA
biobakery3	13	0.1382	0.1123	22.2639	0.2702	0.1814	33.3333	2.8477
biobakery4	13	0.1426	0.1123	22.0538	0.2702	0.1854	33.3333	2.0672
jams	19	0.0486	0.0664	32.3825	0.3691	0.0961	50.0000	50.1032
wgsa2	18	0.0007	0.0740	33.1873	0.3343	0.1114	50.0000	63.0962
woltka	13	0.2100	0.1239	20.0301	0.1949	0.2352	8.3333	2.3966

Bivariate Linear Regression for Sample S1 in Experiment bmock12 (Species at filter threshold 0.1)



Bivariate Linear Regression for Sample S1 in Experiment bmock12 (Species at filter threshold 0.1)



	Diversity	R ²	MAE	AD	1-BC	RMSE	Sens	FPRA
biobakery3	12	0.1356	0.1216	17.5737	0.2702	0.1942	33.3333	0.0000
biobakery4	12	0.1381	0.1216	17.5288	0.2702	0.1970	33.3333	0.0000
jams	15	0.0106	0.0889	29.2969	0.3336	0.1185	41.6667	48.8878
wgsa2	14	0.1657	0.1300	28.6874	0.0899	0.1862	33.3333	80.5083
woltka	12	0.2143	0.1342	12.4164	0.1949	0.2511	8.3333	0.0000