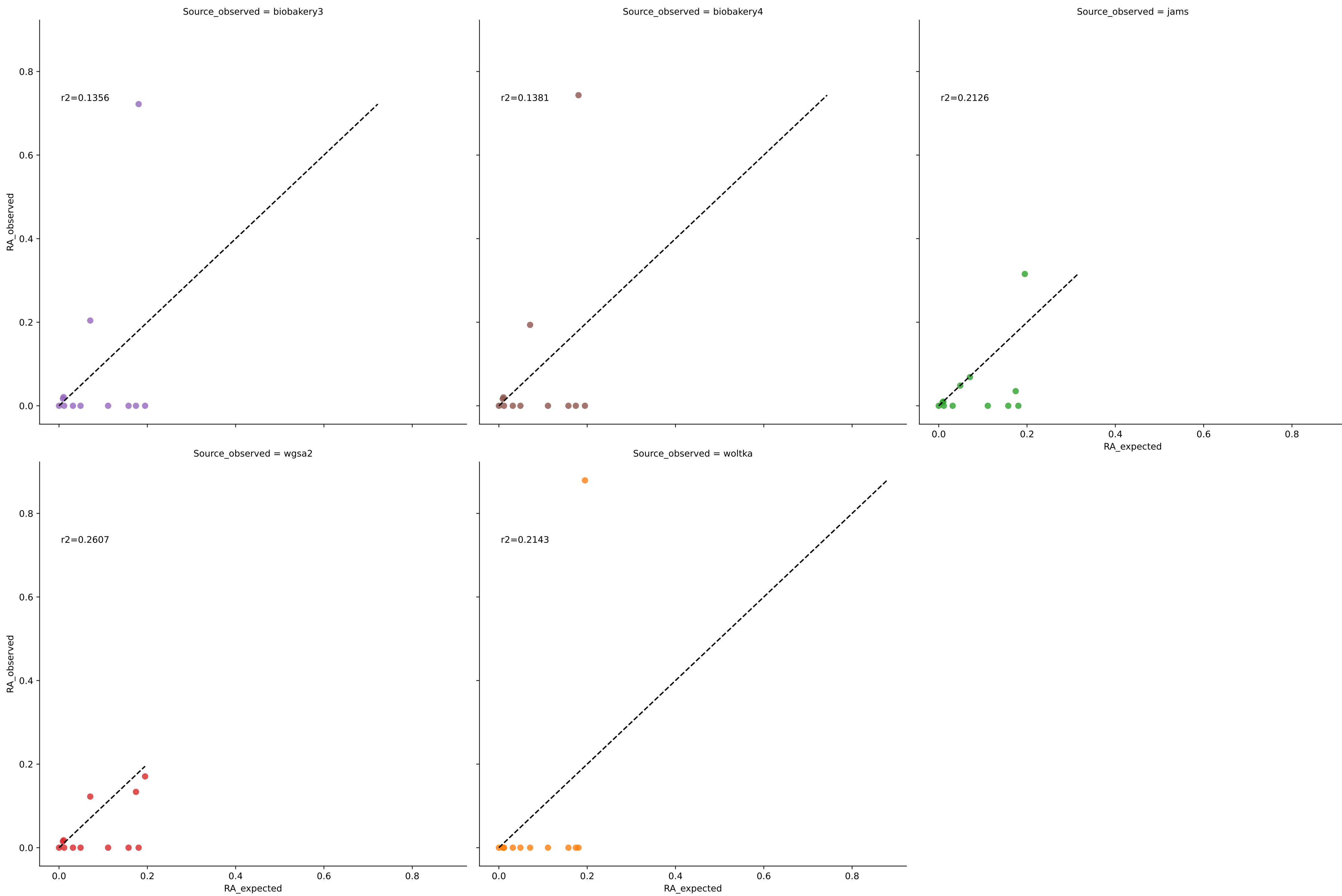
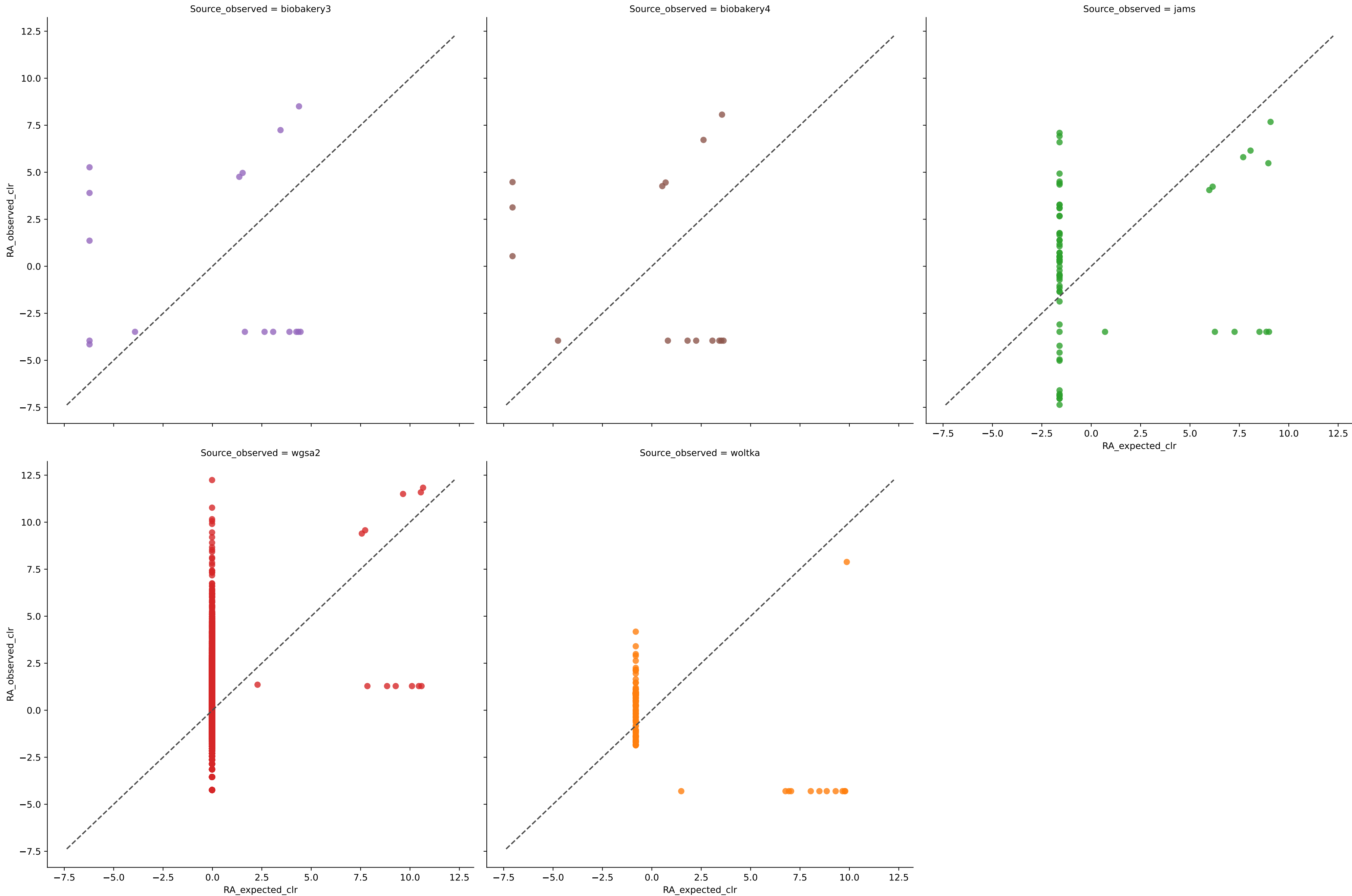


# Bivariate Linear Regression for Sample S1 in Experiment bmock12 (Species at filter threshold 0)

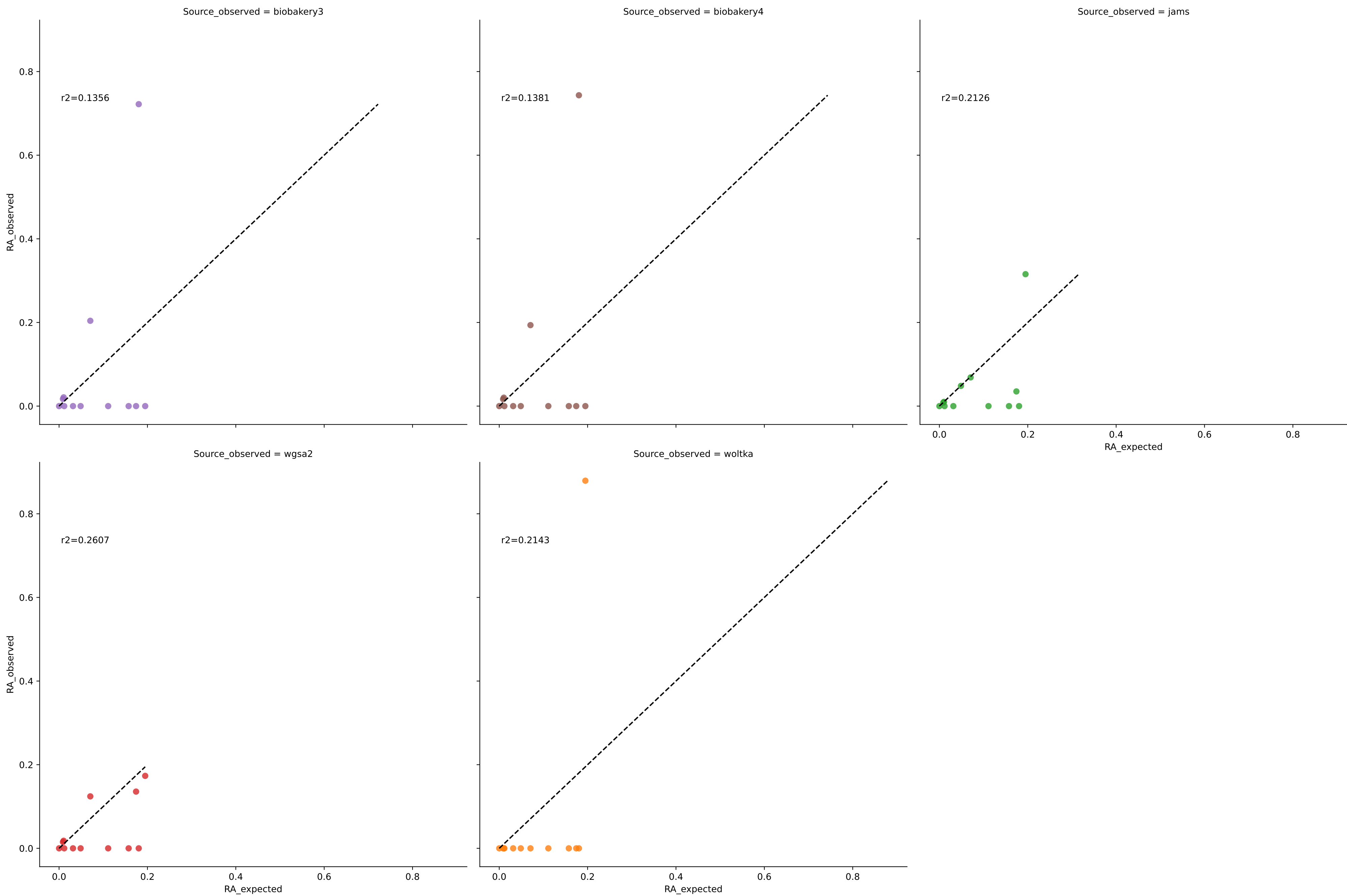


Bivariate Linear Regression for Sample S1 in Experiment bmock12 (Species at filter threshold 0)

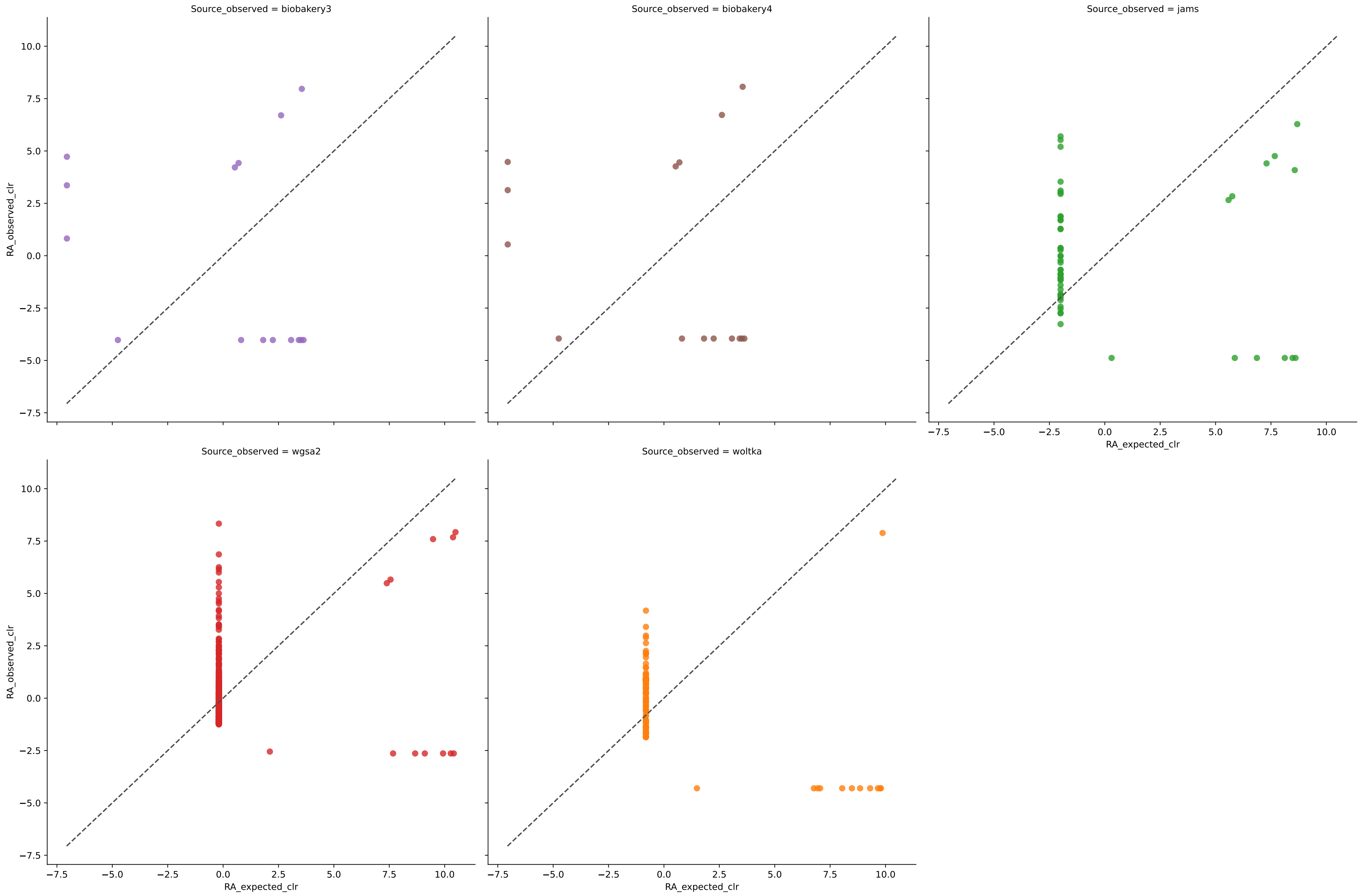


	Diversity	R <sup>2</sup>	MAE	AD	1-BC	RMSE	Sens	FPRA
biobakery3	17	0.1670	0.0859	26.4923	0.2702	0.1574	33.3333	3.6049
biobakery4	15	0.1578	0.0973	25.9132	0.2702	0.1716	33.3333	2.6303
jams	66	0.1721	0.0192	39.5470	0.3655	0.0508	50.0000	51.3979
wgsa2	5630	0.2162	0.0002	177.2455	0.3941	0.0052	50.0000	54.0805
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)

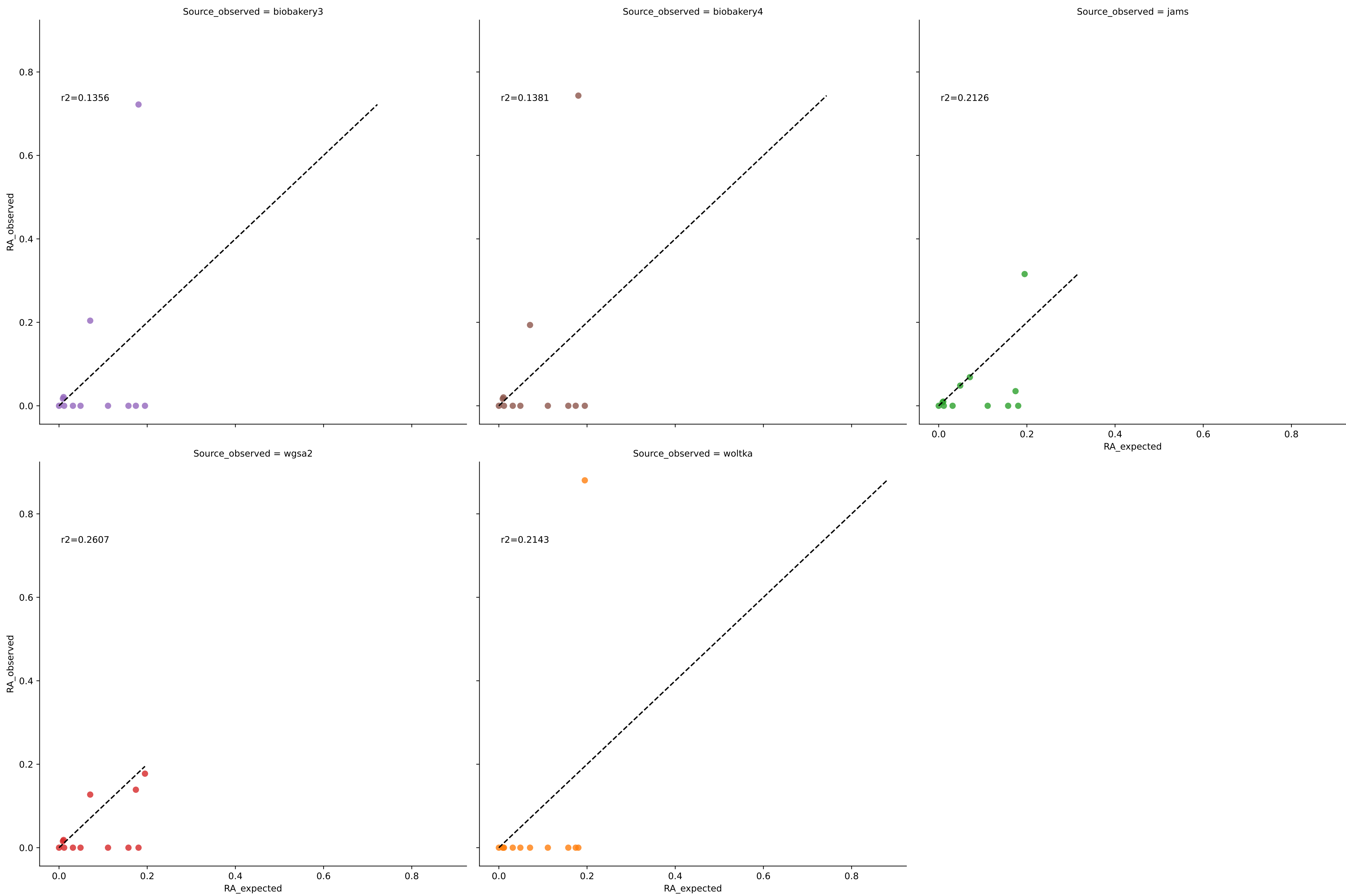


Bivariate Linear Regression for Sample S1 in Experiment bmock12 (Species at filter threshold 1e-05)

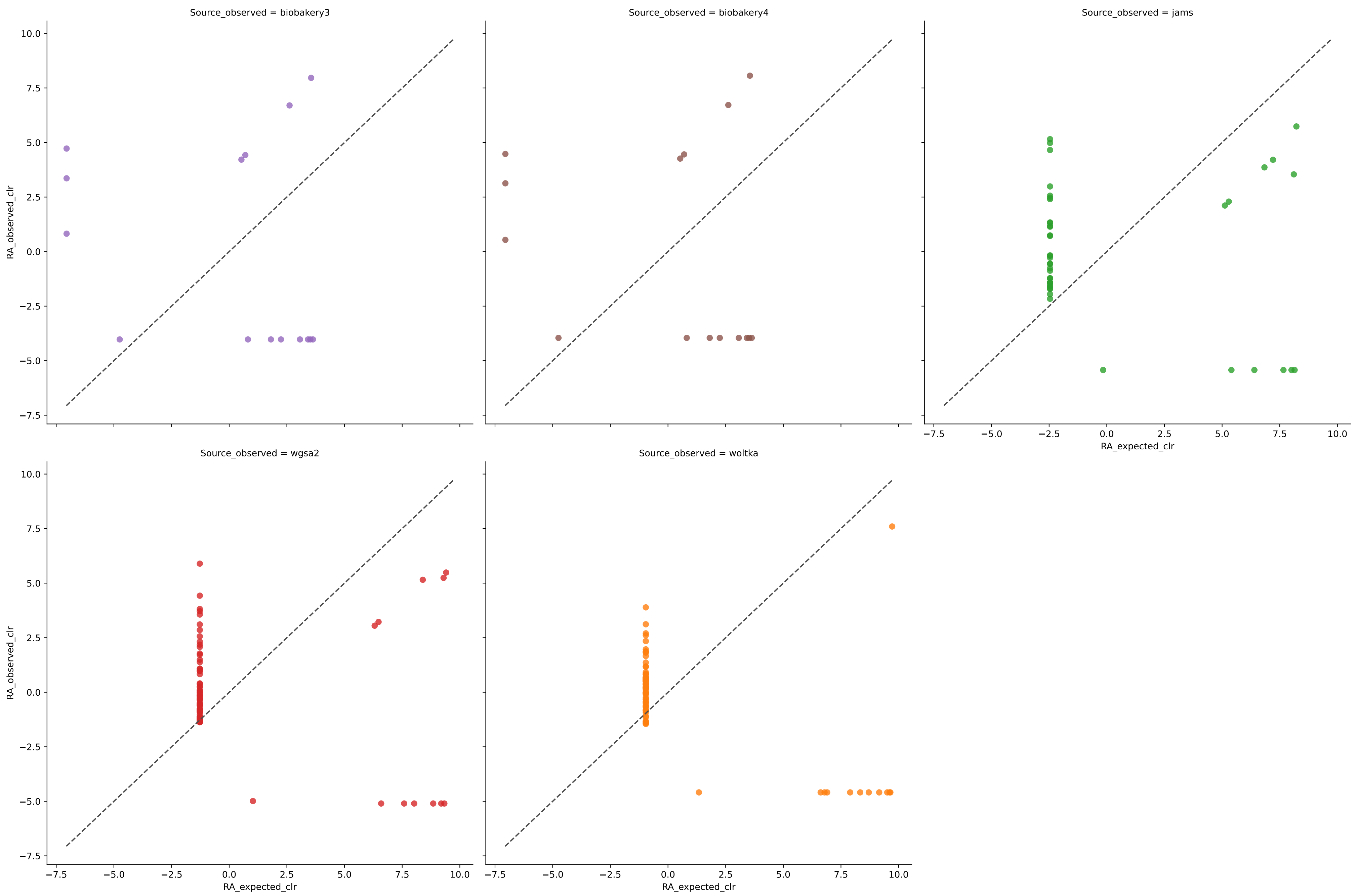


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
biobakery3	15	0.1539	0.0973	26.2902	0.2702	0.1676	33.3333	3.6044
biobakery4	15	0.1578	0.0973	25.9132	0.2702	0.1716	33.3333	2.6303
jams	53	0.1596	0.0239	35.5864	0.3655	0.0566	50.0000	51.3971
wgsa2	550	0.2104	0.0022	43.3137	0.3987	0.0167	50.0000	53.3737
woltka	130	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)

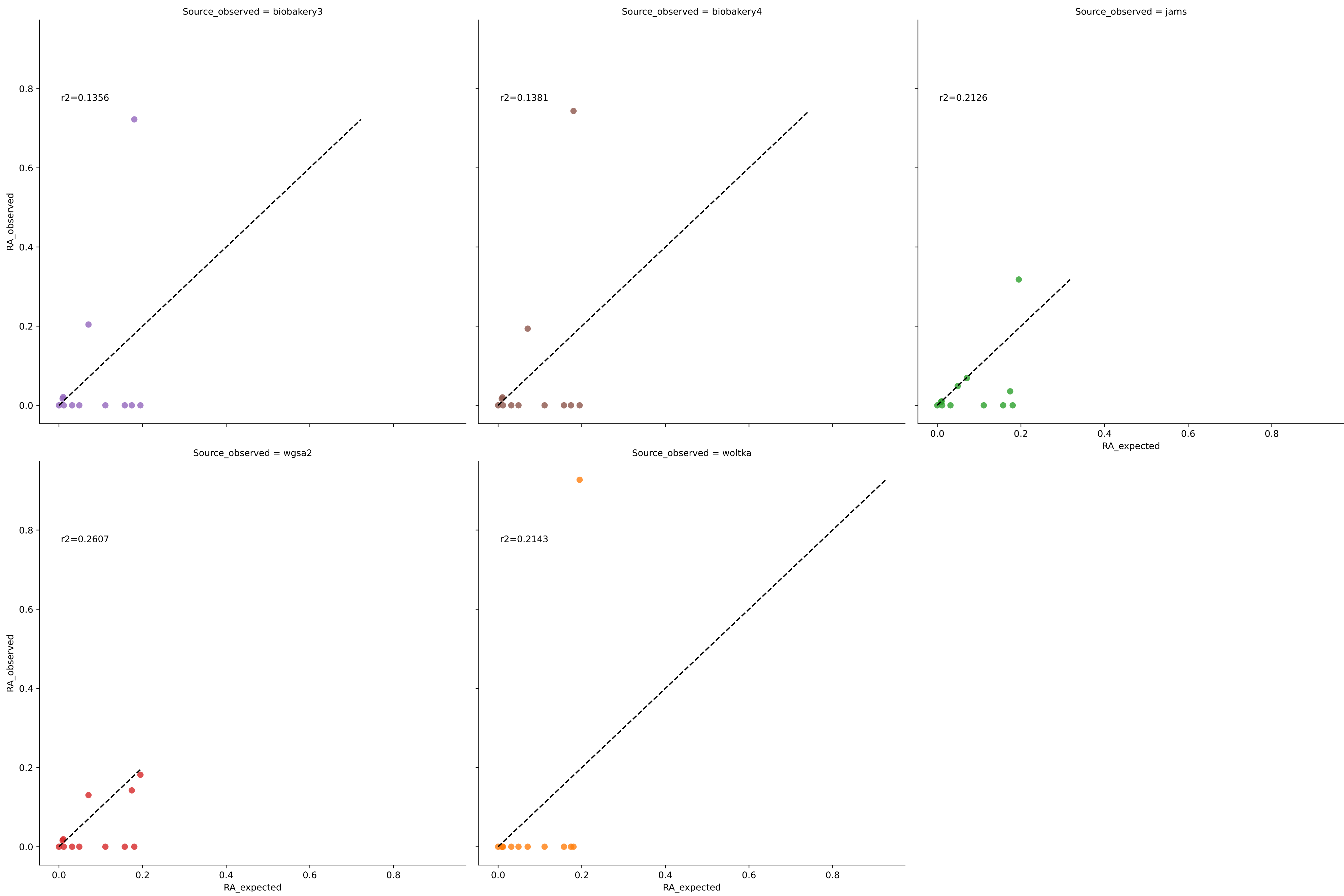


Bivariate Linear Regression for Sample S1 in Experiment bmock12 (Species at filter threshold 0.0001)

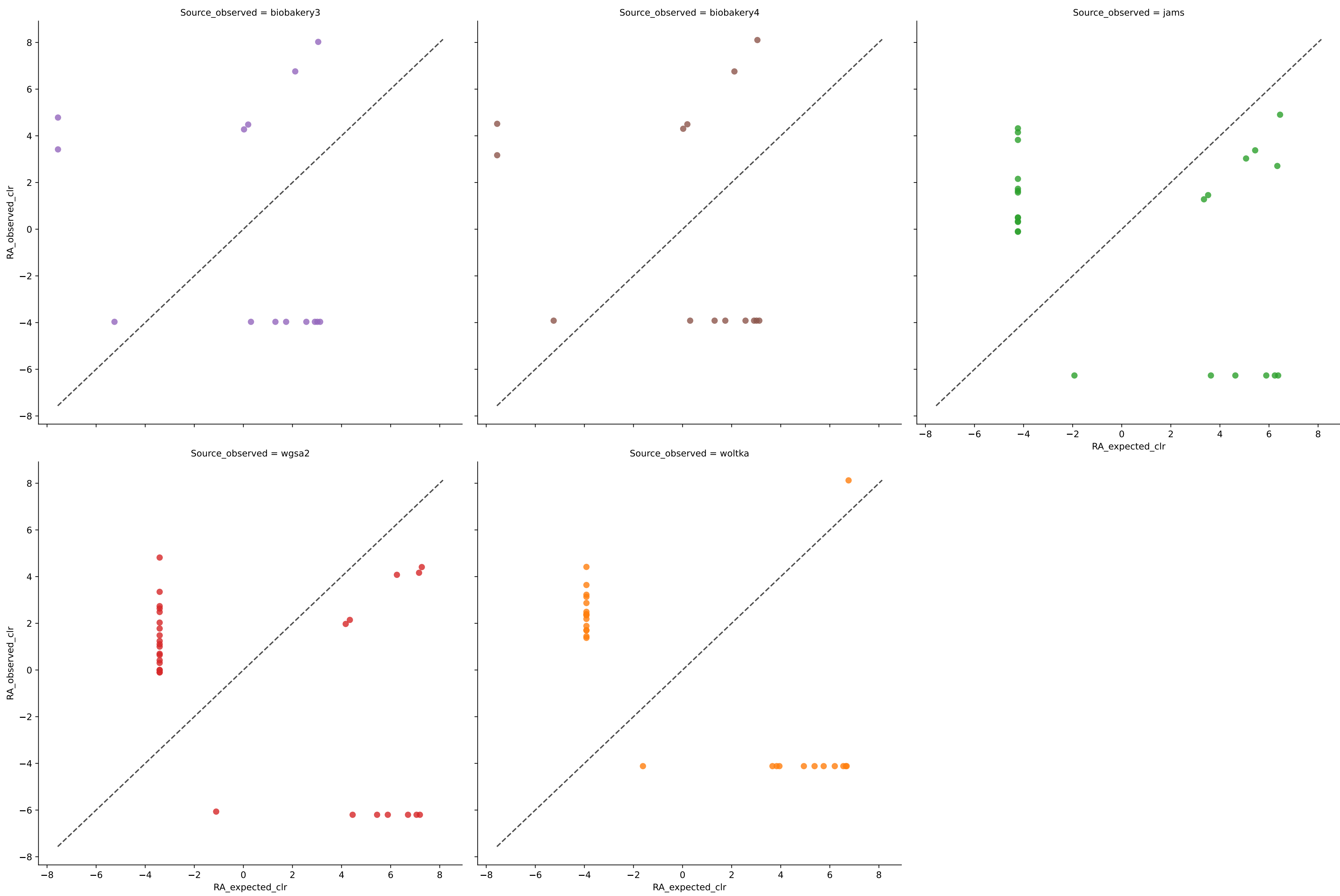


	Diversity	R^2	MAE	AD	1-BC	RMSE	Sens	FPRA
biobakery3	15	0.1539	0.0973	26.2902	0.2702	0.1676	33.3333	3.6044
biobakery4	15	0.1578	0.0973	25.9132	0.2702	0.1716	33.3333	2.6303
jams	43	0.1446	0.0295	35.5374	0.3656	0.0629	50.0000	51.3670
wgsa2	83	0.1744	0.0143	38.9001	0.4062	0.0434	50.0000	52.2500
woltka	110	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)



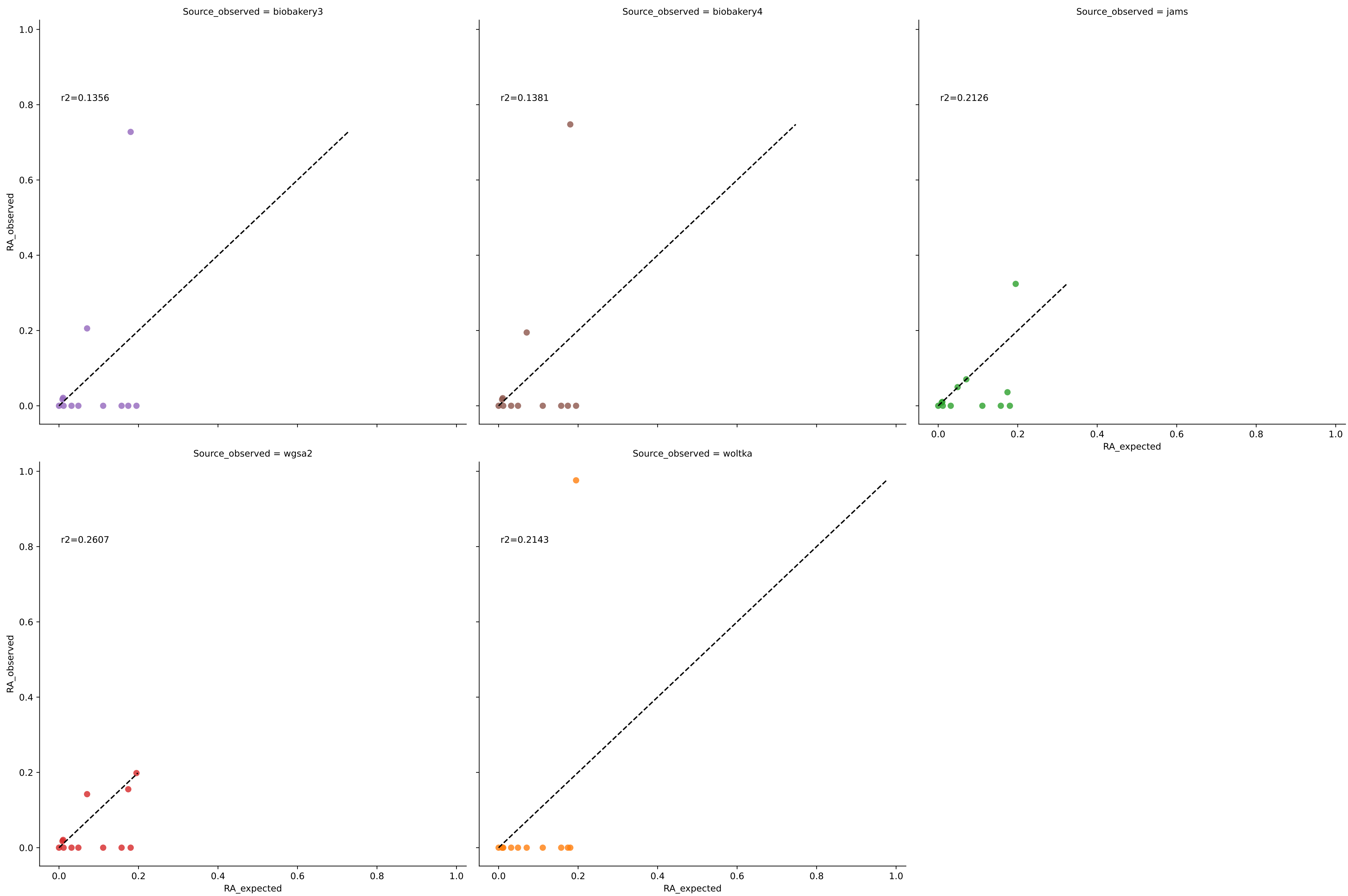
Bivariate Linear Regression for Sample S1 in Experiment bmock12 (Species at filter threshold 0.001)



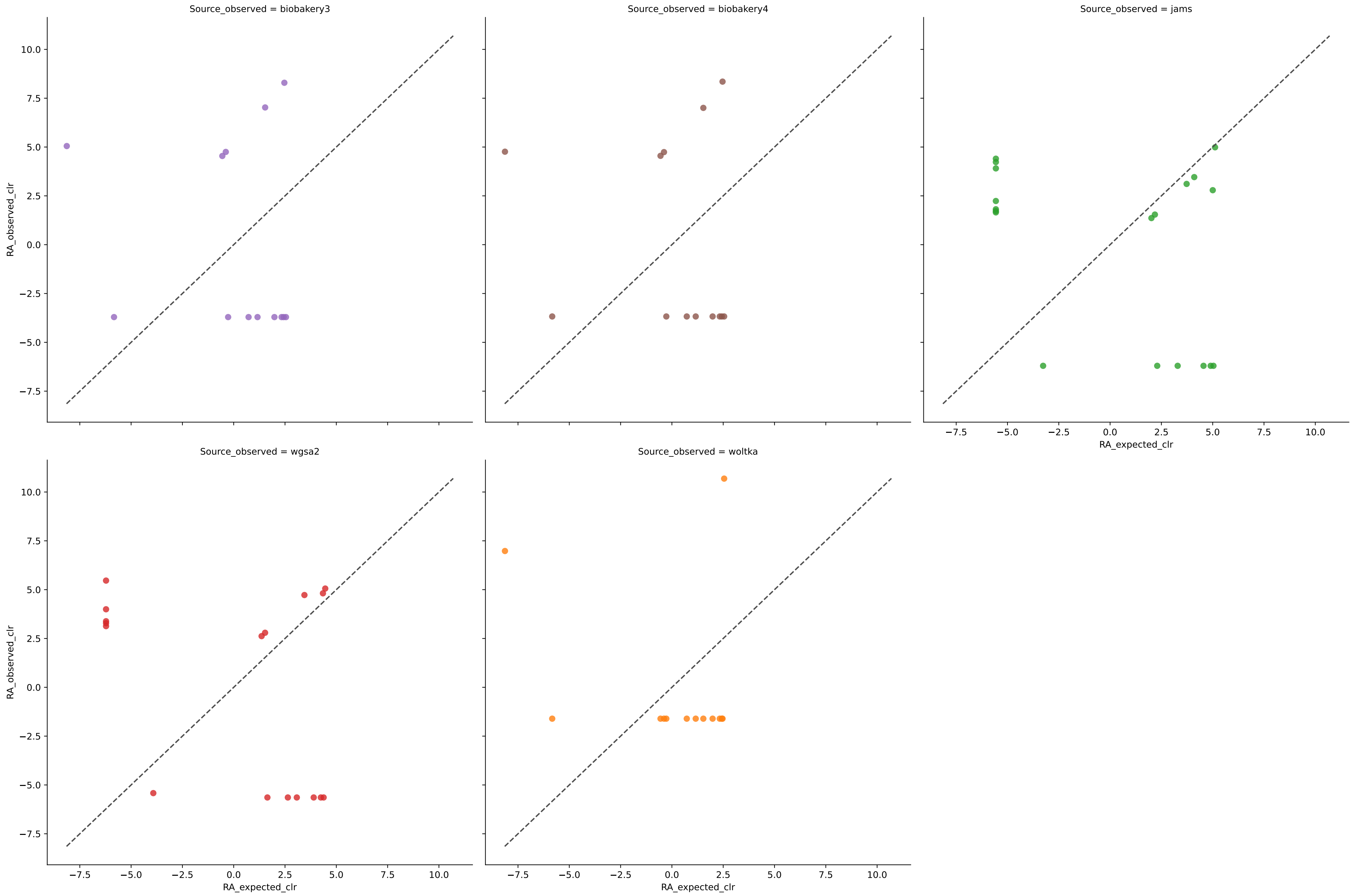
	Diversity	R <sup>2</sup>	MAE	AD	1-BC	RMSE	Sens	FPRA
biobakery3	14	0.1458	0.1043	24.9967	0.2702	0.1736	33.3333	3.5494
biobakery4	14	0.1502	0.1043	24.6946	0.2702	0.1777	33.3333	2.5913
jams	25	0.0872	0.0507	34.7101	0.3668	0.0828	50.0000	51.0282
wgsa2	31	0.1018	0.0378	38.0086	0.4139	0.0719	50.0000	51.0827
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)

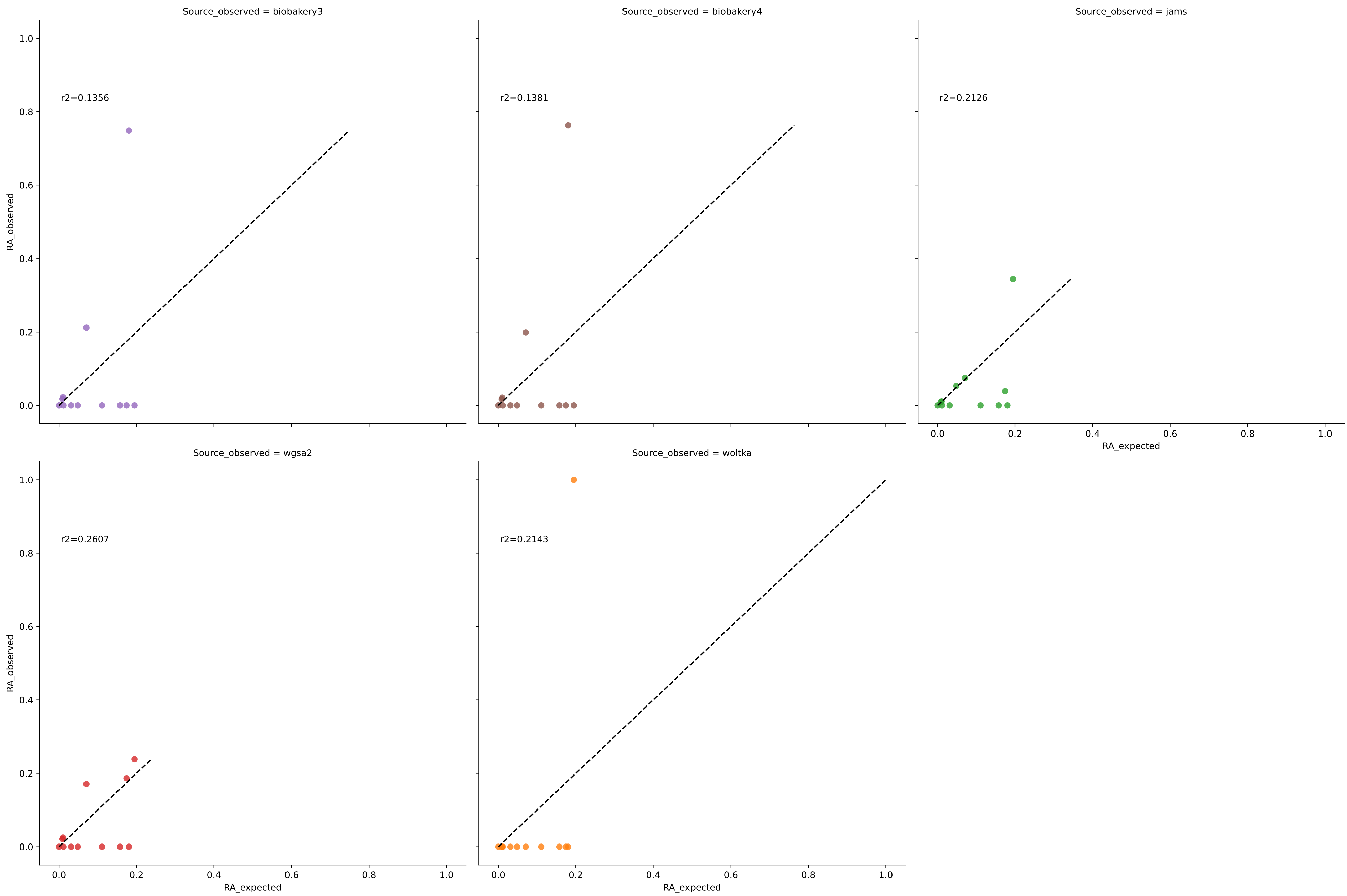


Bivariate Linear Regression for Sample S1 in Experiment bmock12 (Species at filter threshold 0.01)

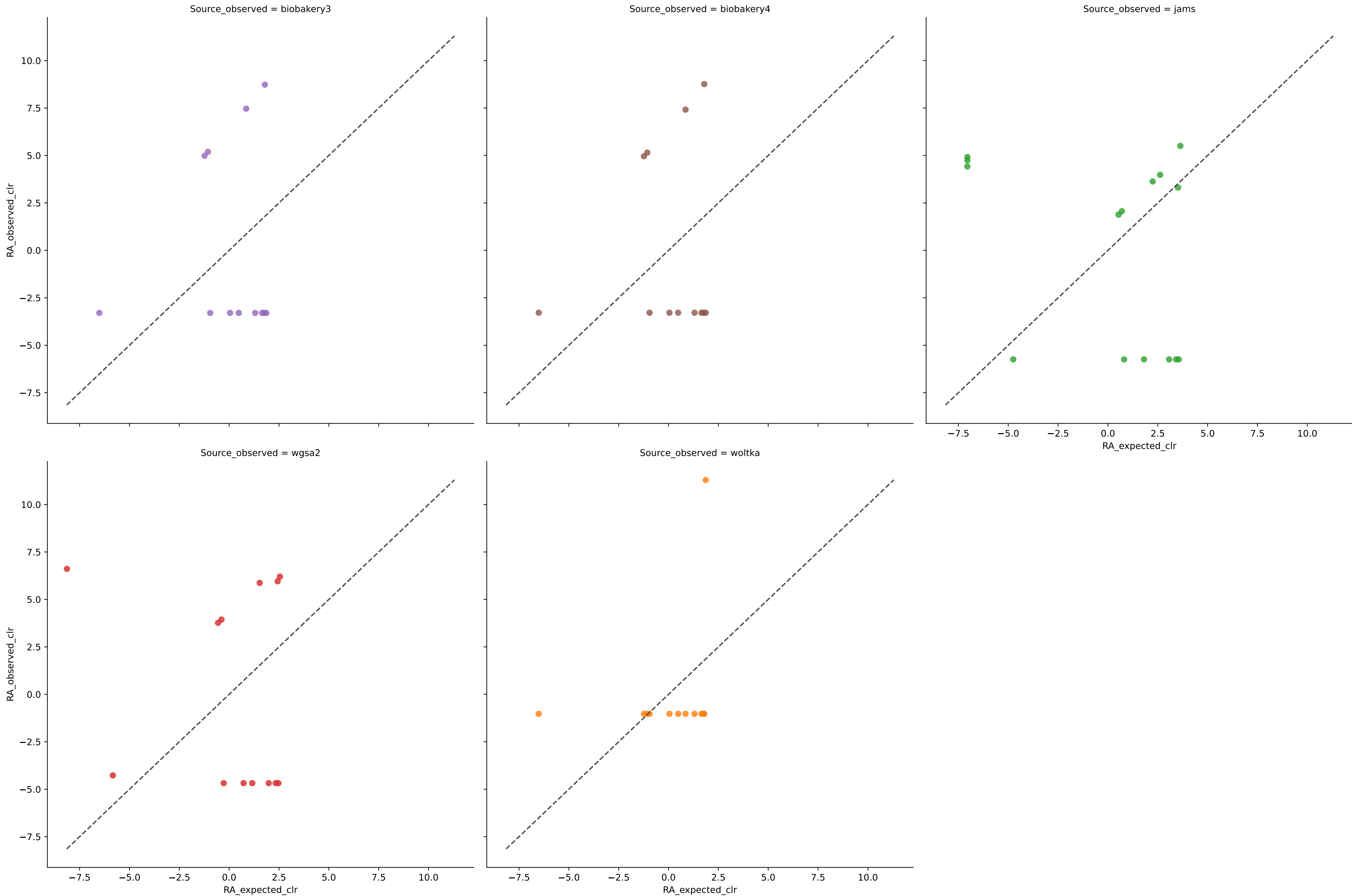


	Diversity	R <sup>2</sup>	MAE	AD	1-BC	RMSE	Sens	FPRA
biobakery3	13	0.1382	0.1123	22.2639	0.2702	0.1814	33.3333	2.8477
biobakery4	13	0.1426	0.1123	22.0538	0.2702	0.1854	33.3333	2.0672
jams	19	0.0486	0.0664	32.3825	0.3691	0.0961	50.0000	50.1032
wgsa2	17	0.0278	0.0659	31.7086	0.4400	0.1017	50.0000	46.6336
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 <sup>2</sup>	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	0.0206	0.0838	27.8266	0.3718	0.1120	50.0000	47.0174
wgsa2	13	0.0176	0.0832	23.0059	0.4591	0.1283	50.0000	35.8338
woltka	12	0.2143	0.1342	12.4164	0.1949	0.2511	8.3333	0.0000