

Co-inoculum of *Beauveria brongniartii* and *B. bassiana* shows in vitro different metabolic behaviour in comparison to single inoculums

Canfora L.¹, Abu Samra N.^{1§}, Tartanus M.², Łabanowska B.H.², Benedetti A.¹, Pinzari F.^{1,3*} and Malusà E.^{2,4}

¹Council for Agricultural Research and Economics, Research Centre for Agriculture and Environment, Via della Navicella 2-4, 00184 Rome, Italy

²Research Institute of Horticulture, Konstytucji 3 Maja 1, 96-100 Skierniewice, Poland

³Department of Life Sciences, Natural History Museum, Cromwell Road, SW7 5BD London, UK

⁴Council for Agricultural Research and Economics, Research Centre for Engineering and Agro-Food Processing, Strada delle Cacce 73, 10135 Torino, Italy

§ current address: Cirad - av. Agropolis - Génétique et innovation variétale (GIV). TA A-108/01 - 34398 Montpellier Cedex 5 – France

*Corresponding author

Supplementary Information:

Tables: S1, S2, S3, S4, S5

Figures: S1, S2, S3 (=a series of 96 plots)

Table S1. One-sided test Hypothesis: CO metabolism larger than BA or BR. Simultaneous Tests for General Linear Hypotheses. OD 490nm.

Linear Hypotheses:					Estimate	Std. Error	t value	Pr(>t)		
B04-Erythritol	CO	-	BA	<= 0	158.45	16.3	9.719	<	1.00E-04	***
	CO	-	BR	<= 0	163.78	16.3	10.046	<	1.00E-04	***
D03-D-Melezitose	CO	-	BA	<= 0	102.76	16.3	6.303	<	1.00E-04	***
	CO	-	BR	<= 0	219.64	16.3	13.473	<	1.00E-04	***
E04-D-Sorbitol	CO	-	BA	<= 0	50.19	16.3	3.079		0.02217	*
	CO	-	BR	<= 0	203.7	16.3	12.495	<	1.00E-04	***
G10-L-Asparagine	CO	-	BA	<= 0	160.13	16.3	9.822	<	1.00E-04	***
	CO	-	BR	<= 0	287.01	16.3	17.605	<	1.00E-04	***
G11-L-Aspartic Acid	CO	-	BA	<= 0	45.8	16.3	2.809		0.04353	*
	CO	-	BR	<= 0	171.88	16.3	10.543	<	1.00E-04	***
G12-Glutamic Acid	CO	-	BA	<= 0	62.31	16.3	3.822		0.00293	**
	CO	-	BR	<= 0	172.8	16.3	10.599	<	1.00E-04	***

Signif. codes: 0 '***', 0.001 '**', 0.01 '*', 0.05 '.', 0.1 ' ', 1. (Adjusted p values reported -- single-step method)

Table S2. One-sided Test. Hypothesis: CO growth larger than that of BA or BR. Simultaneous Tests for General Linear Hypotheses. OD 750nm.

Hypotheses:											
					Estimate	Std. Error	t value	Pr(>t)			
B04-Erythritol	CO	-	BA	<=	0	154.94	12.94	11.975	<	1.00E-04	***
	CO	-	BR	<=	0	154.59	12.94	11.949	<	1.00E-04	***
D03-D-Melezitose	CO	-	BA	<=	0	96.36	12.94	7.448	<	1.00E-04	***
	CO	-	BR	<=	0	204.51	12.94	15.807	<	1.00E-04	***
E04-D-Sorbitol	CO	-	BA	<=	0	45.67	12.94	3.53		0.00665	**
	CO	-	BR	<=	0	187.01	12.94	14.454	<	1.00E-04	***
G10-L-Asparagine	CO	-	BA	<=	0	101.06	12.94	7.811	<	1.00E-04	***
	CO	-	BR	<=	0	153.59	12.94	11.871	<	1.00E-04	***
G11-L-Aspartic Acid	CO	-	BA	<=	0	10.99	12.94	0.85		0.89011	
	CO	-	BR	<=	0	45.05	12.94	3.482		0.00758	**
G12-Glutamic Acid	CO	-	BA	<=	0	26.41	12.94	2.041		0.22892	
	CO	-	BR	<=	0	78.01	12.94	6.03	<	1.00E-04	***

Signif. codes: 0 '***', 0.001 '**', 0.01 '*', 0.05 '.', 0.1 ' ', 1. (Adjusted p values reported -- single-step method)

Table S3. Two-sided Test Hypothesis: CO BA BR different. Simultaneous Tests for General Linear Hypotheses. OD 490nm. Multiple Comparisons of Means: User-defined Contrasts.

Hypotheses:					Estimate	Std. Error	t value	Pr(>t)	
A08-D-Arabinose	BR	-	BA	= 0	28.842	16.369	1.762	0.915	
	CO	-	BA	= 0	-1.925	16.369	-0.118	1.000	
	CO	-	BR	= 0	-30.767	16.369	-1.88	0.856	
A10-D-Arabitol	BR	-	BA	= 0	-75.816	16.369	-4.632	0.001	***
	CO	-	BA	= 0	-11.398	16.369	-0.696	1.000	
	CO	-	BR	= 0	64.418	16.369	3.935	0.007	**
A11-Arbutin	BR	-	BA	= 0	-49.474	16.369	-3.022	0.112	
	CO	-	BA	= 0	-30.813	16.369	-1.882	0.855	
	CO	-	BR	= 0	18.661	16.369	1.14	1.000	
A12-D-Cellobiose	BR	-	BA	= 0	-189.558	16.369	-11.581	< 0.000	***
	CO	-	BA	= 0	-17.52	16.369	-1.07	1.000	
	CO	-	BR	= 0	172.038	16.369	10.51	< 0.000	***
C07-2-Keto-D-Gluconic Acid	BR	-	BA	= 0	122.241	16.369	7.468	< 0.000	***
	CO	-	BA	= 0	49.966	16.369	3.053	0.103	
	CO	-	BR	= 0	-72.275	16.369	-4.415	0.001	**
C08- α -D-Lactose	BR	-	BA	= 0	-73.72	16.369	-4.504	0.001	***
	CO	-	BA	= 0	12.939	16.369	0.79	1.000	
	CO	-	BR	= 0	86.659	16.369	5.294	< 0.000	***
D08- β -Methyl-D-Glucoside	BR	-	BA	= 0	-163.698	16.369	-10.001	< 0.000	***
	CO	-	BA	= 0	-10.301	16.369	-0.629	1.000	
	CO	-	BR	= 0	153.397	16.369	9.371	< 0.000	***
E06-Stachynose	BR	-	BA	= 0	-155.953	16.369	-9.528	< 0.000	***
	CO	-	BA	= 0	2.725	16.369	0.166	1.000	
	CO	-	BR	= 0	158.679	16.369	9.694	< 0.000	***
F06-p-Hydroxyphenyl-acetic acid	BR	-	BA	= 0	15.838	16.369	0.968	1.000	
	CO	-	BA	= 0	-3.81	16.369	-0.233	1.000	
	CO	-	BR	= 0	-19.648	16.369	-1.2	0.999	
G06-N-Acetyly-L-Glutamic Acid	BR	-	BA	= 0	87.302	16.369	5.333	< 0.000	***
	CO	-	BA	= 0	41.233	16.369	2.519	0.364	
	CO	-	BR	= 0	-46.069	16.369	-2.814	0.190	
H03-L-Phenylalanine	BR	-	BA	= 0	-64.796	16.369	-3.959	0.006	**
	CO	-	BA	= 0	-33.269	16.369	-2.032	0.754	
	CO	-	BR	= 0	31.526	16.369	1.926	0.828	
H10-Adenosine	BR	-	BA	= 0	17.833	16.369	1.089	1.000	
	CO	-	BA	= 0	-1.154	16.369	-0.071	1.000	
	CO	-	BR	= 0	-18.988	16.369	-1.16	1.000	
H11-Uridine	BR	-	BA	= 0	33.579	16.369	2.051	0.739	
	CO	-	BA	= 0	3.126	16.369	0.191	1.000	
	CO	-	BR	= 0	-30.453	16.369	-1.86	0.867	
A08-D-Arabinose	BR	-	BA	= 0	28.842	16.369	1.762	0.915	

CO	-	BA	=	0	-1.925	16.369	-0.118	1.000
CO	-	BR	=	0	-30.767	16.369	-1.88	0.856

Signif. codes: 0 '***', 0.001 '**', 0.01 '*', 0.05 '.', 0.1 ' ', 1. (Adjusted p values reported -- single-step method)

Table S4. Two-sided Test Hypothesis: CO BA BR different. Simultaneous Tests for General Linear Hypotheses. OD 750nm. Multiple Comparisons of Means: User-defined Contrasts.

		Hypotheses:			
		Estimate	Std. Error	t value	Pr(>t)
A08-D-Arabinose	BR - BA = 0	18.935	13.186	1.436	0.991
	CO - BA = 0	-1.284	13.186	-0.097	1.000
	CO - BR = 0	-20.218	13.186	-1.533	0.979
A10-D-Arabitol	BR - BA = 0	-45.445	13.186	-3.446	0.033 *
	CO - BA = 0	-0.878	13.186	-0.067	1.000
	CO - BR = 0	44.567	13.186	3.380	0.040 *
A11-Arbutin	BR - BA = 0	-7.273	13.186	-0.552	1.000
	CO - BA = 0	-6.353	13.186	-0.482	1.000
	CO - BR = 0	0.920	13.186	0.070	1.000
A12-D-Cellobiose	BR - BA = 0	-186.252	13.186	-14.125	< 0.000 ***
	CO - BA = 0	-12.156	13.186	-0.922	1.000
	CO - BR = 0	174.097	13.186	13.203	< 0.000 ***
C07-2-Keto-D-Gluconic Acid	BR - BA = 0	16.649	13.186	1.263	0.998
	CO - BA = 0	28.605	13.186	2.169	0.645
	CO - BR = 0	11.956	13.186	0.907	1.000
C08- α -D-Lactose	BR - BA = 0	-70.300	13.186	-5.331	< 0.000 ***
	CO - BA = 0	14.923	13.186	1.132	1.000
	CO - BR = 0	85.223	13.186	6.463	< 0.000 ***
D08- β -Methyl-D-Glucoside	BR - BA = 0	-157.446	13.186	-11.940	< 0.000 ***
	CO - BA = 0	-8.189	13.186	-0.621	1.000
	CO - BR = 0	149.257	13.186	11.319	< 0.000 ***
E06-Stachynose	BR - BA = 0	-145.328	13.186	-11.021	< 0.000 ***
	CO - BA = 0	13.852	13.186	1.051	1.000
	CO - BR = 0	159.181	13.186	12.072	< 0.000 ***
F06-p-Hydroxyphenyl-acetic acid	BR - BA = 0	25.955	13.186	1.968	0.800
	CO - BA = 0	-0.715	13.186	-0.054	1.000
	CO - BR = 0	-26.670	13.186	-2.023	0.761
G06-N-Acetyly-L-Glutamic Acid	BR - BA = 0	47.140	13.186	3.575	0.022 *
	CO - BA = 0	22.603	13.186	1.714	0.934
	CO - BR = 0	-24.537	13.186	-1.861	0.867
H03-L-Phenylalanine	BR - BA = 0	4.115	13.186	0.312	1.000
	CO - BA = 0	-2.061	13.186	-0.156	1.000
	CO - BR = 0	-6.176	13.186	-0.468	1.000
H10-Adenosine	BR - BA = 0	20.661	13.186	1.567	0.973
	CO - BA = 0	-3.877	13.186	-0.294	1.000
	CO - BR = 0	-24.538	13.186	-1.861	0.867
H11-Uridine	BR - BA = 0	23.678	13.186	1.796	0.900
	CO - BA = 0	0.393	13.186	0.030	1.000
	CO - BR = 0	-23.285	13.186	-1.766	0.914
A08-D-Arabinose	BR - BA = 0	18.935	13.186	1.436	0.991

CO - BA = 0	-1.284	13.186	-0.097	1.000
CO - BR = 0	-20.218	13.186	-1.533	0.979

Signif. codes: 0 '***', 0.001 '**', 0.01 '*', 0.05 '.', 0.1 ' ', 1. (Adjusted p values reported -- single-step method)

Table S5. Kendall's tau coefficient used to measure the ordinal association between respiration (OD 490) and growth (OD 750 nm) data. The colour gradient is used in the table to graphically represent the degree of correlation between growth and respiration (green=low degree, red=high degree).

Substrate	tau CO	2-sided p-value	tau BA	2-sided p-value	tau BR	2-sided p-value
A.Cyclodextrin	0.695	<2.22e-16	0.788	<2.22e-16	0.639	<2.22e-16
a.D.Glucose	0.886	<2.22e-16	0.949	<2.22e-16	0.968	<2.22e-16
a.D.Lactose	0.98	<2.22e-16	0.976	<2.22e-16	0.896	<2.22e-16
A.Keto.glutaricAcid	0.868	<2.22e-16	0.846	<2.22e-16	0.912	<2.22e-16
a.Methyl.D.Galactoside	0.769	<2.22e-16	0.826	<2.22e-16	0.575	3.58E-07
a.Methyl.D.Glucoside	0.729	<2.22e-16	0.743	<2.22e-16	0.537	1.67E-06
Adenosine	0.681	<2.22e-16	0.741	<2.22e-16	0.293	0.0089771
Adenosine.5.Monophosphate	0.783	<2.22e-16	0.756	<2.22e-16	0.692	<2.22e-16
Adonitol	0.894	<2.22e-16	0.959	<2.22e-16	0.895	<2.22e-16
Alaninamide	0.729	<2.22e-16	0.751	<2.22e-16	0.64	<2.22e-16
Amygdalin	0.816	<2.22e-16	0.87	<2.22e-16	0.957	<2.22e-16
Arbutin	0.795	<2.22e-16	0.861	<2.22e-16	0.978	<2.22e-16
b.Cyclodextrin	0.756	<2.22e-16	0.786	<2.22e-16	0.61	<2.22e-16
B.Hydroxy.butyricAcid	0.825	<2.22e-16	0.792	<2.22e-16	0.765	<2.22e-16
b.Methyl.D.Galactoside	0.828	<2.22e-16	0.828	<2.22e-16	0.861	<2.22e-16
b.Methyl.D.Glucoside	0.884	<2.22e-16	0.93	<2.22e-16	0.978	<2.22e-16
BromosuccinicAcid	0.822	<2.22e-16	0.772	<2.22e-16	0.885	<2.22e-16
D.Arabinose	0.773	<2.22e-16	0.797	<2.22e-16	0.872	<2.22e-16
D.Arabitol	0.975	<2.22e-16	0.972	<2.22e-16	0.946	<2.22e-16
D.Cellobiose	0.874	<2.22e-16	0.87	<2.22e-16	0.968	<2.22e-16
D.Fructose	0.962	<2.22e-16	0.97	<2.22e-16	0.949	<2.22e-16
D.Galactose	0.967	<2.22e-16	0.975	<2.22e-16	0.982	<2.22e-16
D.GalacturonicAcid	0.835	<2.22e-16	0.834	<2.22e-16	0.83	<2.22e-16
D.GluconicAcid	0.837	<2.22e-16	0.862	<2.22e-16	0.866	<2.22e-16
D.Glucosamine	0.933	<2.22e-16	0.934	<2.22e-16	0.894	<2.22e-16
D.GlucuronicAcid	0.802	<2.22e-16	0.746	<2.22e-16	0.606	1.19E-07
D.LacticAcidMethylEster	0.662	<2.22e-16	0.685	<2.22e-16	0.618	<2.22e-16
D.MalicAcid	0.959	<2.22e-16	0.914	<2.22e-16	0.854	<2.22e-16
D.Mannitol	0.866	<2.22e-16	0.866	<2.22e-16	0.916	<2.22e-16
D.MannitolBA	0.916	<2.22e-16	0.973	<2.22e-16	0.973	<2.22e-16
D.Mannose	0.791	<2.22e-16	0.897	<2.22e-16	0.981	<2.22e-16
D.Melezitose	0.957	<2.22e-16	0.954	<2.22e-16	0.507	5.96E-06
D.Melibiose	0.97	<2.22e-16	0.98	<2.22e-16	0.583	2.38E-07
D.Psicose	0.816	<2.22e-16	0.816	<2.22e-16	0.723	<2.22e-16
D.Raffinose	0.727	<2.22e-16	0.894	<2.22e-16	0.968	<2.22e-16
D.Ribose	0.951	<2.22e-16	0.964	<2.22e-16	0.948	<2.22e-16
D.SaccharinAcid	0.616	<2.22e-16	0.636	<2.22e-16	0.471	2.68E-05
D.Sorbitol	0.808	<2.22e-16	0.889	<2.22e-16	0.922	<2.22e-16
D.Tagatose	0.709	<2.22e-16	0.691	<2.22e-16	0.623	<2.22e-16
D.Trehalose	0.826	<2.22e-16	0.914	<2.22e-16	0.976	<2.22e-16
D.Xylose	0.947	<2.22e-16	0.949	<2.22e-16	0.943	<2.22e-16

Dextrin	0.913	<2.22e-16	0.928	<2.22e-16	0.941	<2.22e-16
FumaricAcid	0.972	<2.22e-16	0.868	<2.22e-16	0.887	<2.22e-16
G.Amino.butyricAcid	0.811	<2.22e-16	0.826	<2.22e-16	0.897	<2.22e-16
G.Hydroxy.butyricAcid	0.773	<2.22e-16	0.773	<2.22e-16	0.654	<2.22e-16
Gentibiose	0.727	<2.22e-16	0.88	<2.22e-16	0.989	<2.22e-16
Glucose.1.Phophate	0.951	<2.22e-16	0.959	<2.22e-16	0.897	<2.22e-16
Glucuronamide	0.684	<2.22e-16	0.687	<2.22e-16	0.67	<2.22e-16
Glycerol	0.836	<2.22e-16	0.803	<2.22e-16	0.964	<2.22e-16
Glycogen	0.96	<2.22e-16	0.976	<2.22e-16	0.943	<2.22e-16
Glycyl.L.GlutamicAcid	0.892	<2.22e-16	0.93	<2.22e-16	0.927	<2.22e-16
I.Erythritol	0.968	<2.22e-16	0.921	<2.22e-16	0.943	<2.22e-16
L.Alanine	0.895	<2.22e-16	0.9	<2.22e-16	0.943	<2.22e-16
L.Alanyl.Glycine	0.957	<2.22e-16	0.959	<2.22e-16	0.935	<2.22e-16
L.Arabinose	0.984	<2.22e-16	0.981	<2.22e-16	0.911	<2.22e-16
L.Asparagine	0.938	<2.22e-16	0.943	<2.22e-16	0.875	<2.22e-16
L.AsparticAcid	0.899	<2.22e-16	0.938	<2.22e-16	0.924	<2.22e-16
L.Fucose	0.749	<2.22e-16	0.673	<2.22e-16	0.841	<2.22e-16
L.GlutamicAcid	0.96	<2.22e-16	0.959	<2.22e-16	0.953	<2.22e-16
L.LacticAcid	0.735	<2.22e-16	0.735	<2.22e-16	0.674	<2.22e-16
L.LacticAcidBA	0.674	<2.22e-16	0.613	<2.22e-16	0.613	<2.22e-16
L.MalicAcid	0.9	<2.22e-16	0.868	<2.22e-16	0.903	<2.22e-16
L.Ornithine	0.924	<2.22e-16	0.783	<2.22e-16	0.791	<2.22e-16
L.Phenylalanine	0.787	<2.22e-16	0.856	<2.22e-16	0.538	1.55E-06
L.Proline	0.935	<2.22e-16	0.907	<2.22e-16	0.926	<2.22e-16
L.PyroglutamicAcid	0.897	<2.22e-16	0.842	<2.22e-16	0.932	<2.22e-16
L.Rhamnose	0.738	<2.22e-16	0.754	<2.22e-16	0.538	1.55E-06
L.Serine	0.9	<2.22e-16	0.903	<2.22e-16	0.903	<2.22e-16
L.Sorbose	0.8	<2.22e-16	0.937	<2.22e-16	0.691	<2.22e-16
L.Threonine	0.747	<2.22e-16	0.727	<2.22e-16	0.689	<2.22e-16
Lactulose	0.914	<2.22e-16	0.9	<2.22e-16	0.808	<2.22e-16
m.Inositol	0.927	<2.22e-16	0.935	<2.22e-16	0.935	<2.22e-16
Maltitol	0.809	<2.22e-16	0.793	<2.22e-16	0.716	<2.22e-16
Maltose	0.906	<2.22e-16	0.929	<2.22e-16	0.97	<2.22e-16
Maltotriose	0.867	<2.22e-16	0.875	<2.22e-16	0.974	<2.22e-16
N.Acetyl.D.Galactosamine	0.818	<2.22e-16	0.791	<2.22e-16	0.452	5.58E-05
N.Acetyl.D.Glucosamine	0.967	<2.22e-16	0.967	<2.22e-16	0.972	<2.22e-16
N.Acetyl.D.Mannosamine	0.793	<2.22e-16	0.743	<2.22e-16	0.5	8.46E-06
N.Acetyly.L.GlutamicAcid	0.914	<2.22e-16	0.827	<2.22e-16	0.957	<2.22e-16
P.Hydroxyphenyl.aceticAcid	0.91	<2.22e-16	0.781	<2.22e-16	0.904	<2.22e-16
Palatinose	0.88	<2.22e-16	0.878	<2.22e-16	0.727	<2.22e-16
Putrescine	0.943	<2.22e-16	0.903	<2.22e-16	0.903	<2.22e-16
QuinicAcid	0.773	<2.22e-16	0.71	<2.22e-16	0.511	4.77E-06
Salicin	0.678	<2.22e-16	0.777	<2.22e-16	0.862	<2.22e-16
SebacicAcid	0.67	<2.22e-16	0.723	<2.22e-16	0.922	<2.22e-16
Sedoheptulosan	0.748	<2.22e-16	0.74	<2.22e-16	0.544	1.19E-06
Stachynose	0.891	<2.22e-16	0.87	<2.22e-16	0.883	<2.22e-16
SuccinamicAcid	0.774	<2.22e-16	0.78	<2.22e-16	0.811	<2.22e-16

SuccinicAcid	0.792	<2.22e-16	0.819	<2.22e-16	0.935	<2.22e-16
SuccinicAcidMono.MethylEster	0.838	<2.22e-16	0.636	<2.22e-16	0.738	<2.22e-16
Sucrose	0.527	2.50E-06	0.575	2.38E-07	0.968	<2.22e-16
Turanose	0.86	<2.22e-16	0.895	<2.22e-16	0.921	<2.22e-16
Tween80	0.897	<2.22e-16	0.776	<2.22e-16	0.887	<2.22e-16
Uridine	0.725	<2.22e-16	0.619	<2.22e-16	0.78	<2.22e-16
Water	0.728	<2.22e-16	0.783	<2.22e-16	0.543	1.31E-06
2.AminoEthanol	0.953	<2.22e-16	0.951	<2.22e-16	0.813	<2.22e-16
2.Keto.D.GluconicAcid	0.946	<2.22e-16	0.962	<2.22e-16	0.876	<2.22e-16
Xylitol	0.97	<2.22e-16	0.943	<2.22e-16	0.969	<2.22e-16

Figure S1a. (supplementary material) Average values (n=3) of optical density units (OD) at 490nm (respiration). Phenotype Microarray averaged curves of CO, BA and BR on all the 95 substrates + water. The x-axis shows the measurement time in hours, the y-axis the measured colour intensities in optical density units (OD).

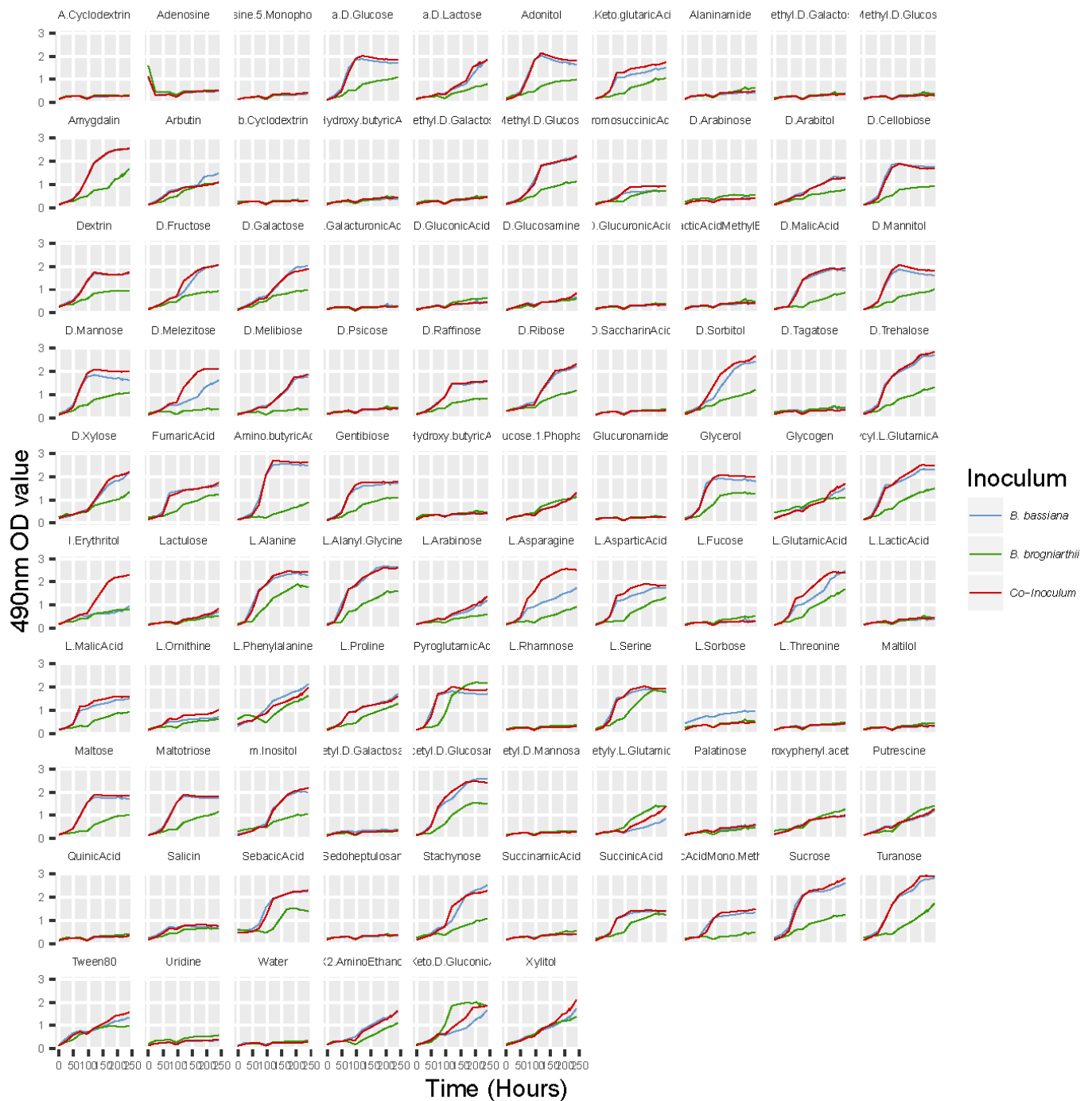
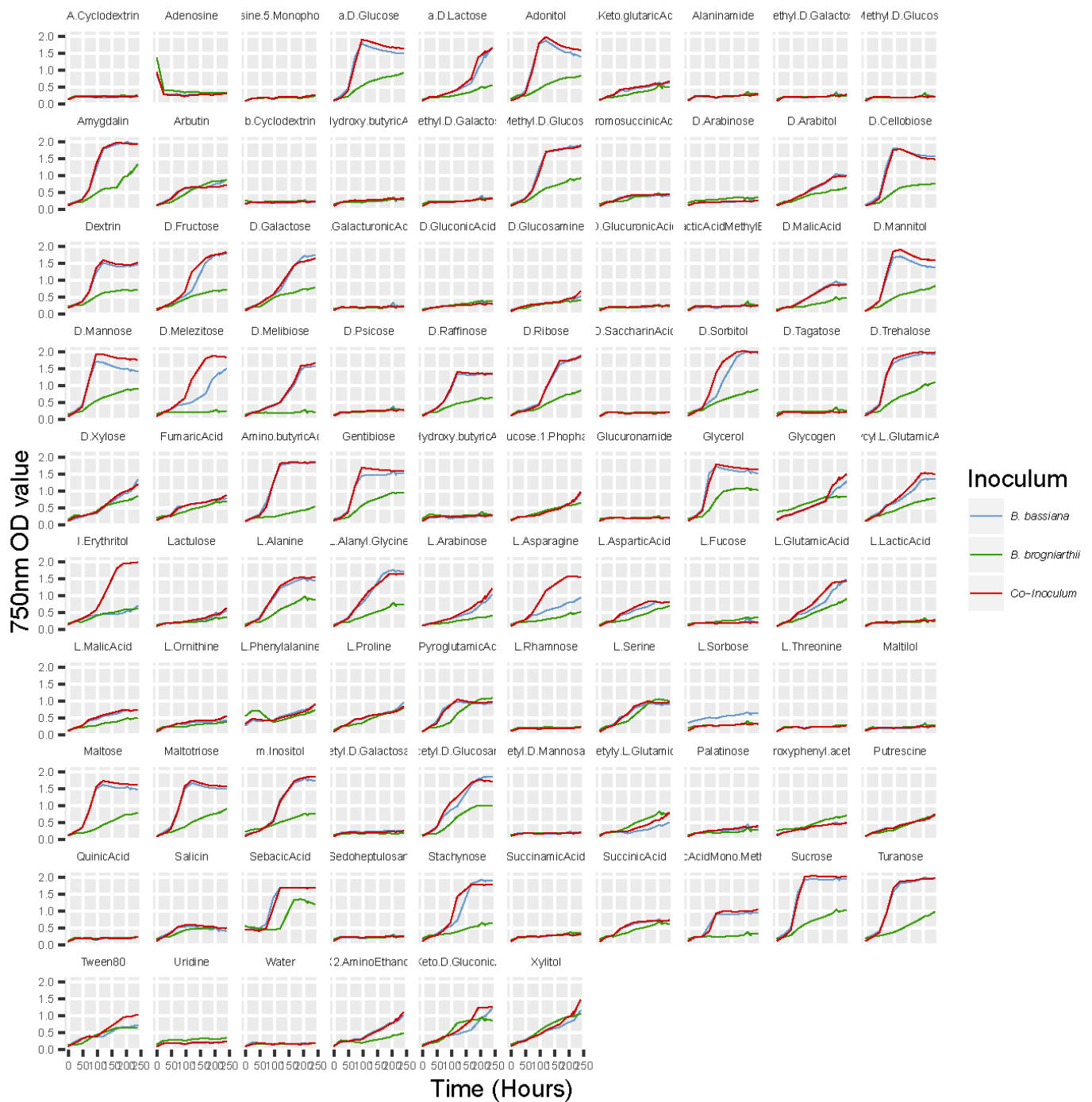
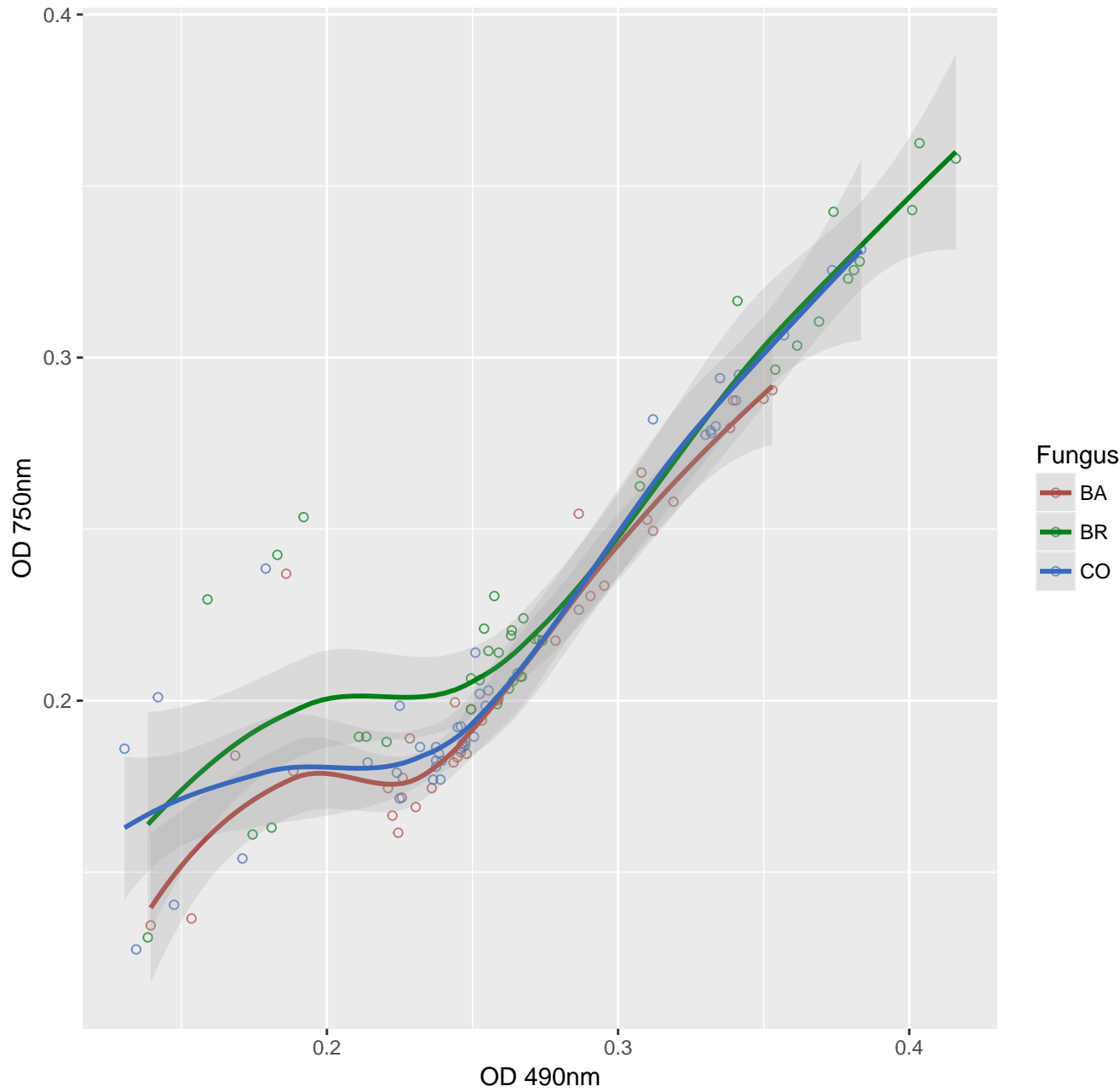


Figure S1b. (supplementary material) Average values (n=3) of optical density units (OD) at 750nm (growth/biomass). Phenotype Microarray averaged curves of CO, BA and BR on all the 95 substrates + water. The x-axis shows the measurement time in hours, the y-axis the measured colour intensities in optical density units (OD).

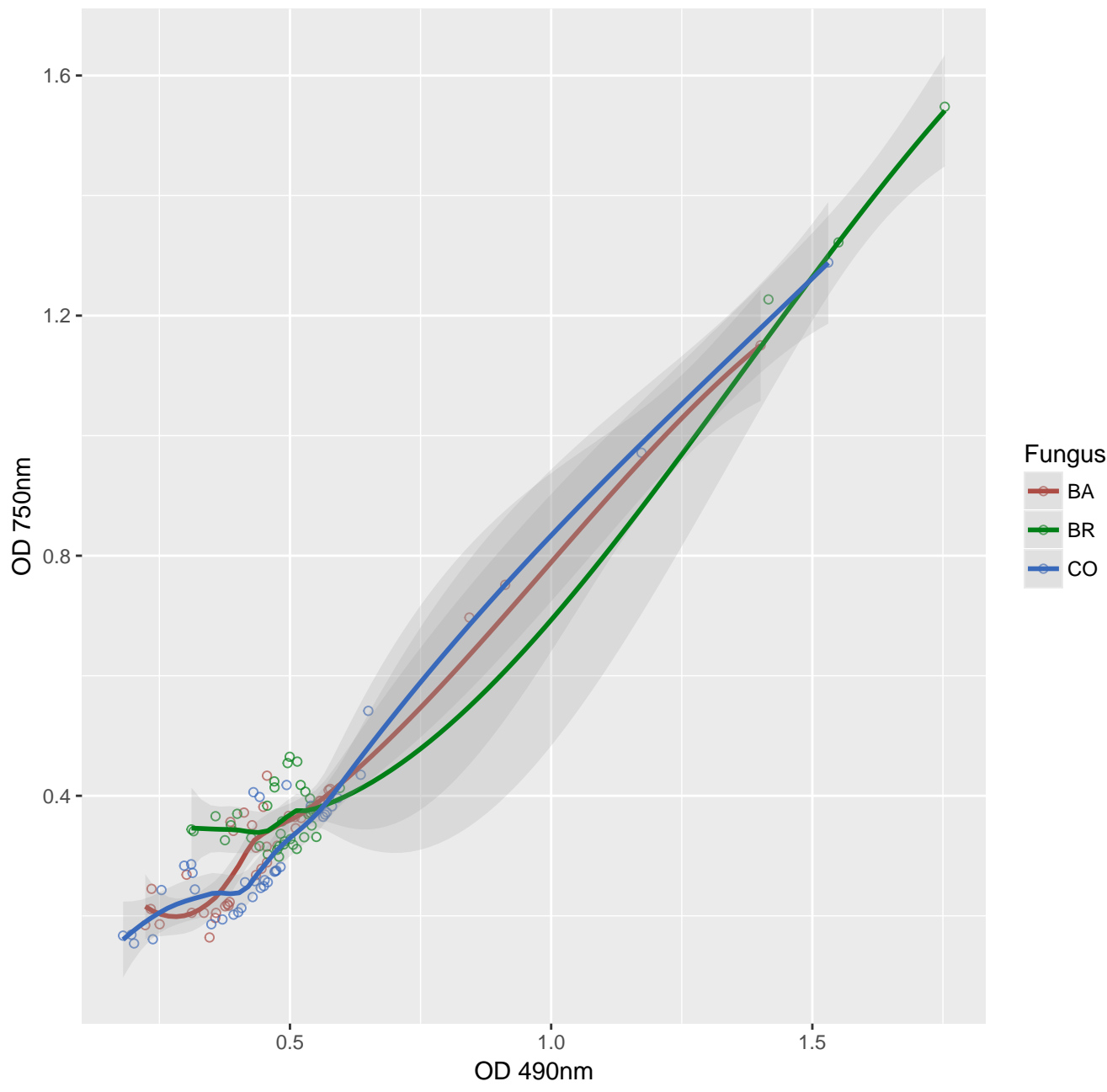


Figures S2 (1 to 96) LOESS (locally weighted scatterplot smoothing) non-parametric regression curves obtained plotting respiration to growth values in time, from all the 96 substrates by the three inoculums.

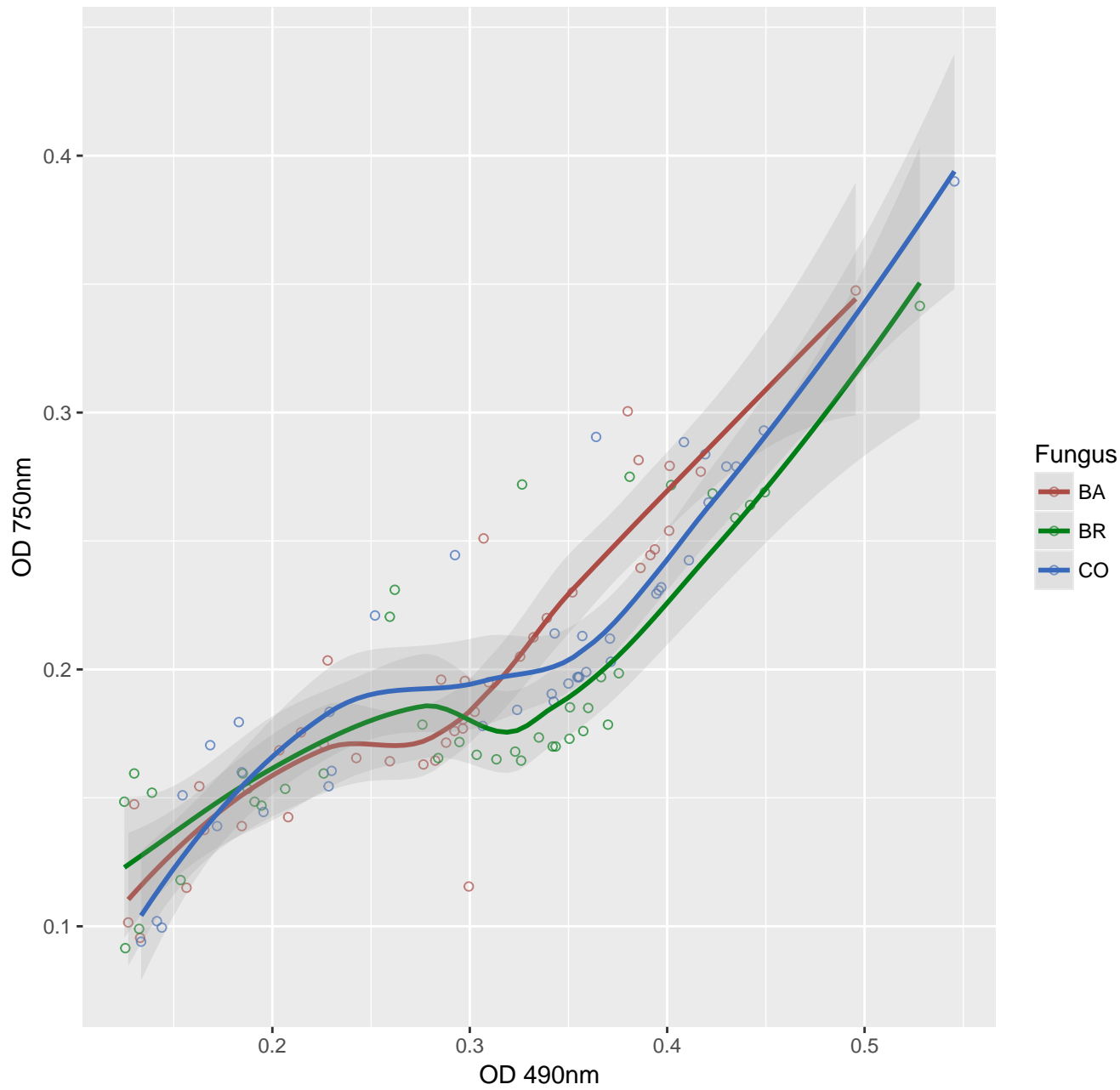
A. Cyclodextrin



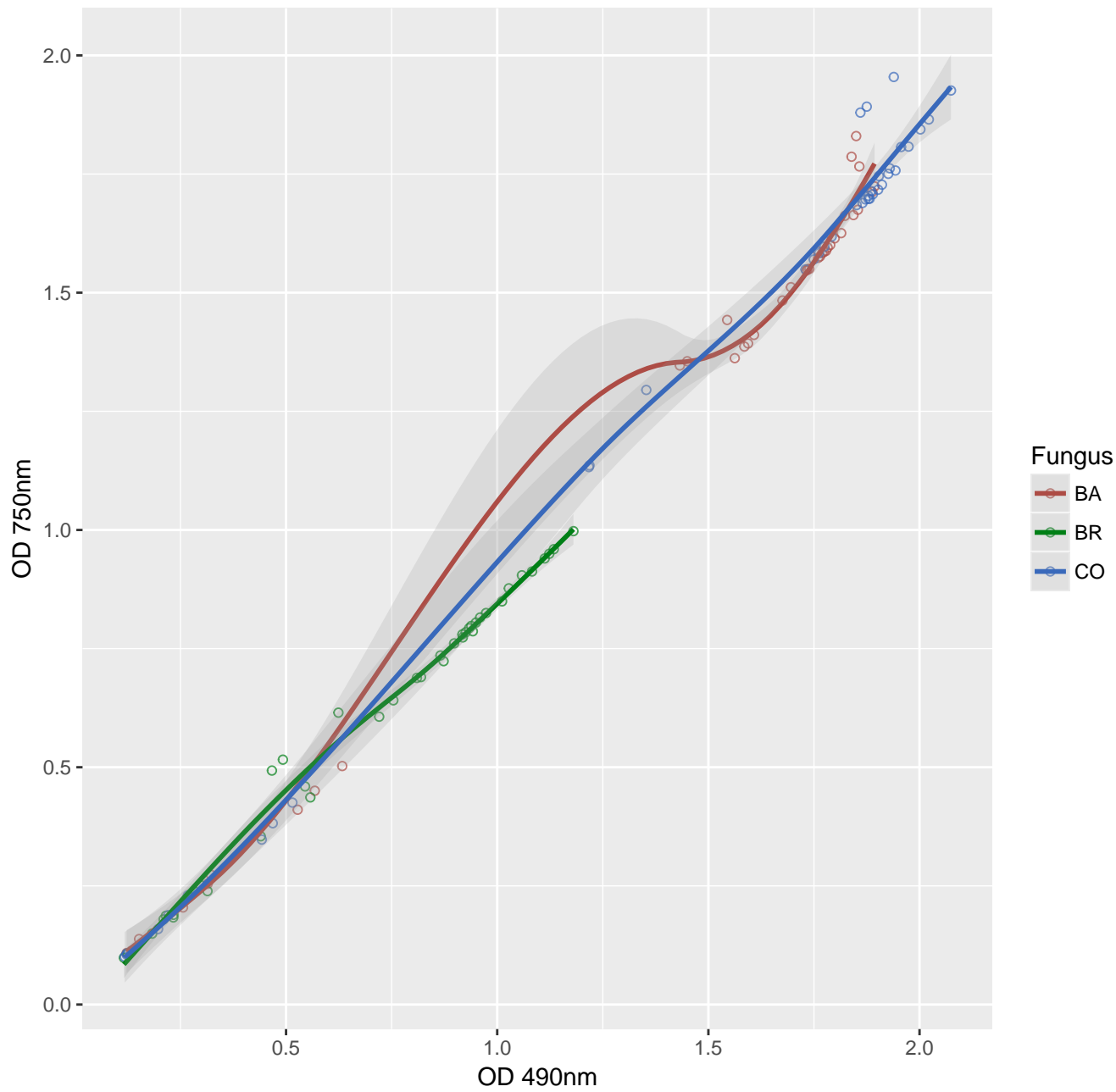
Adenosine



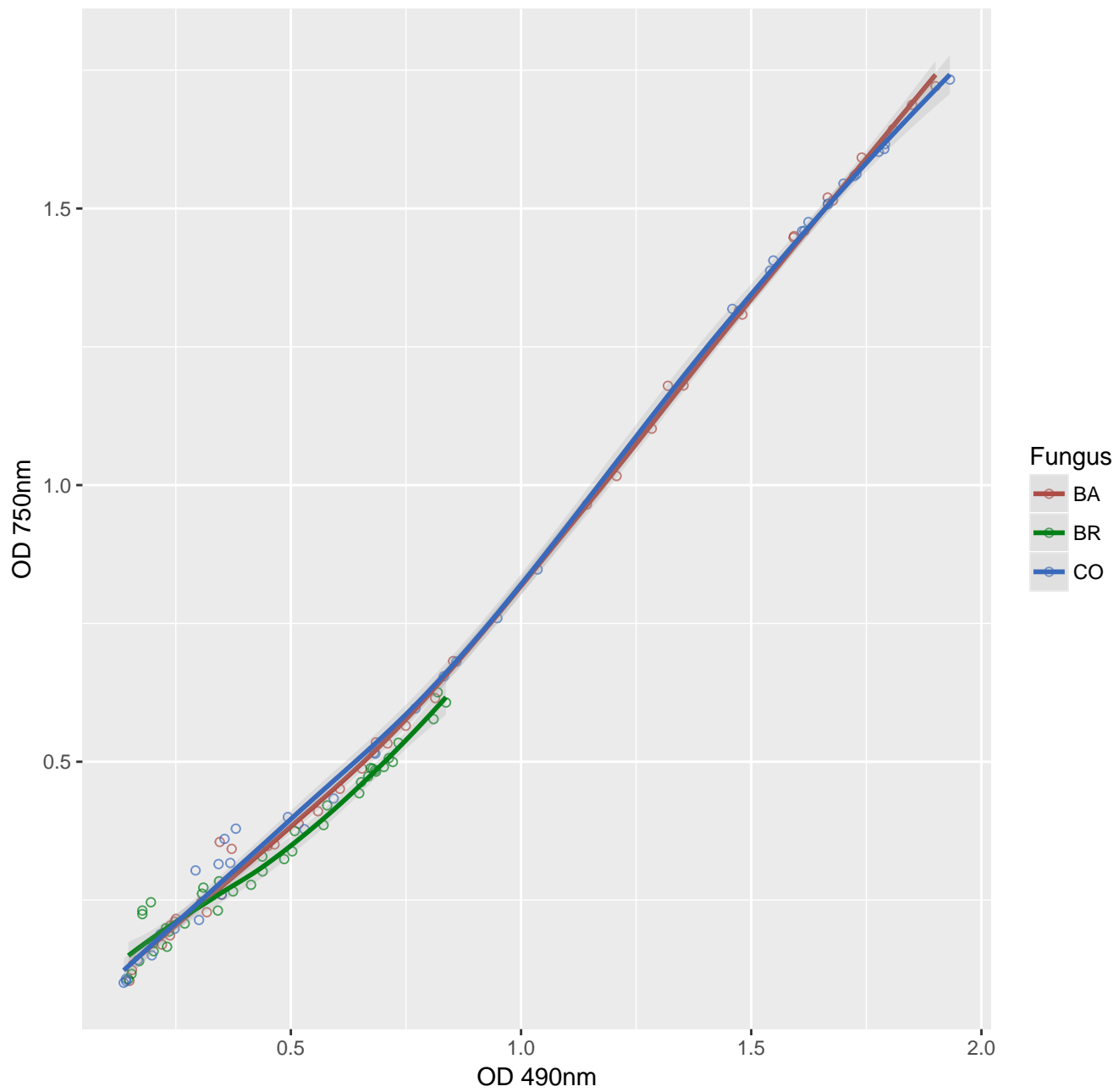
Adenosine.5.Monophosphate



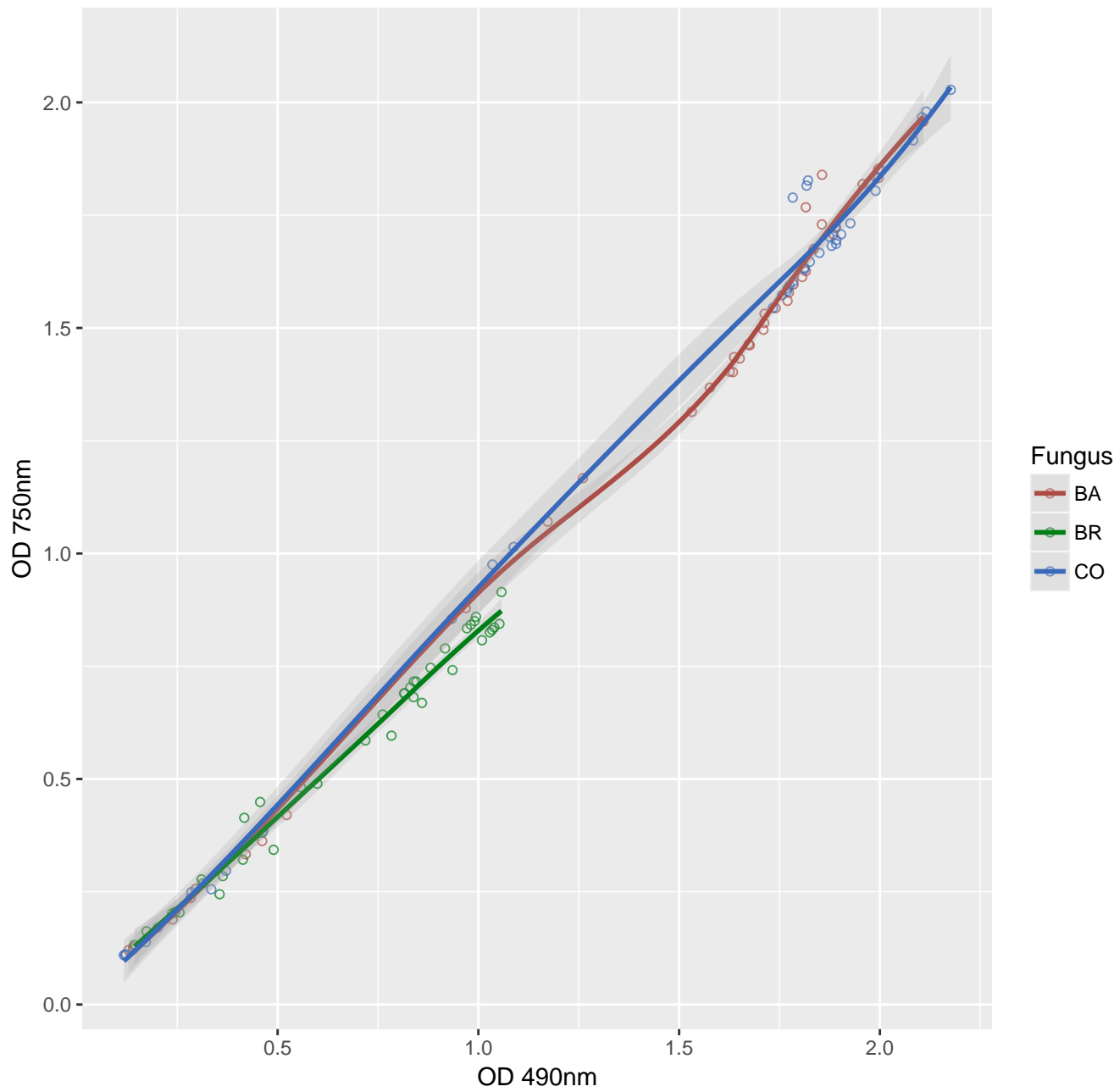
a.D.Glucose



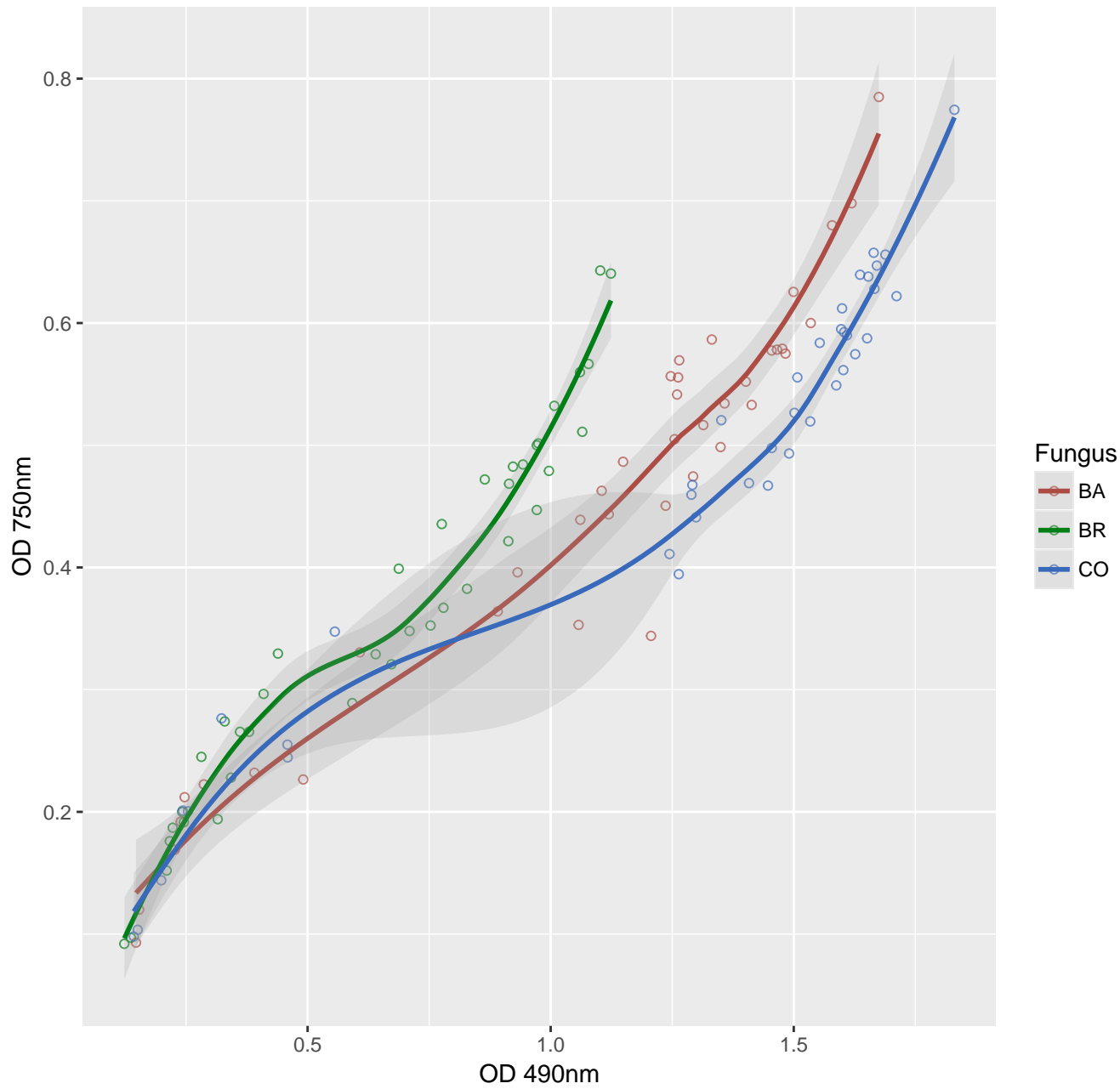
a.D.Lactose



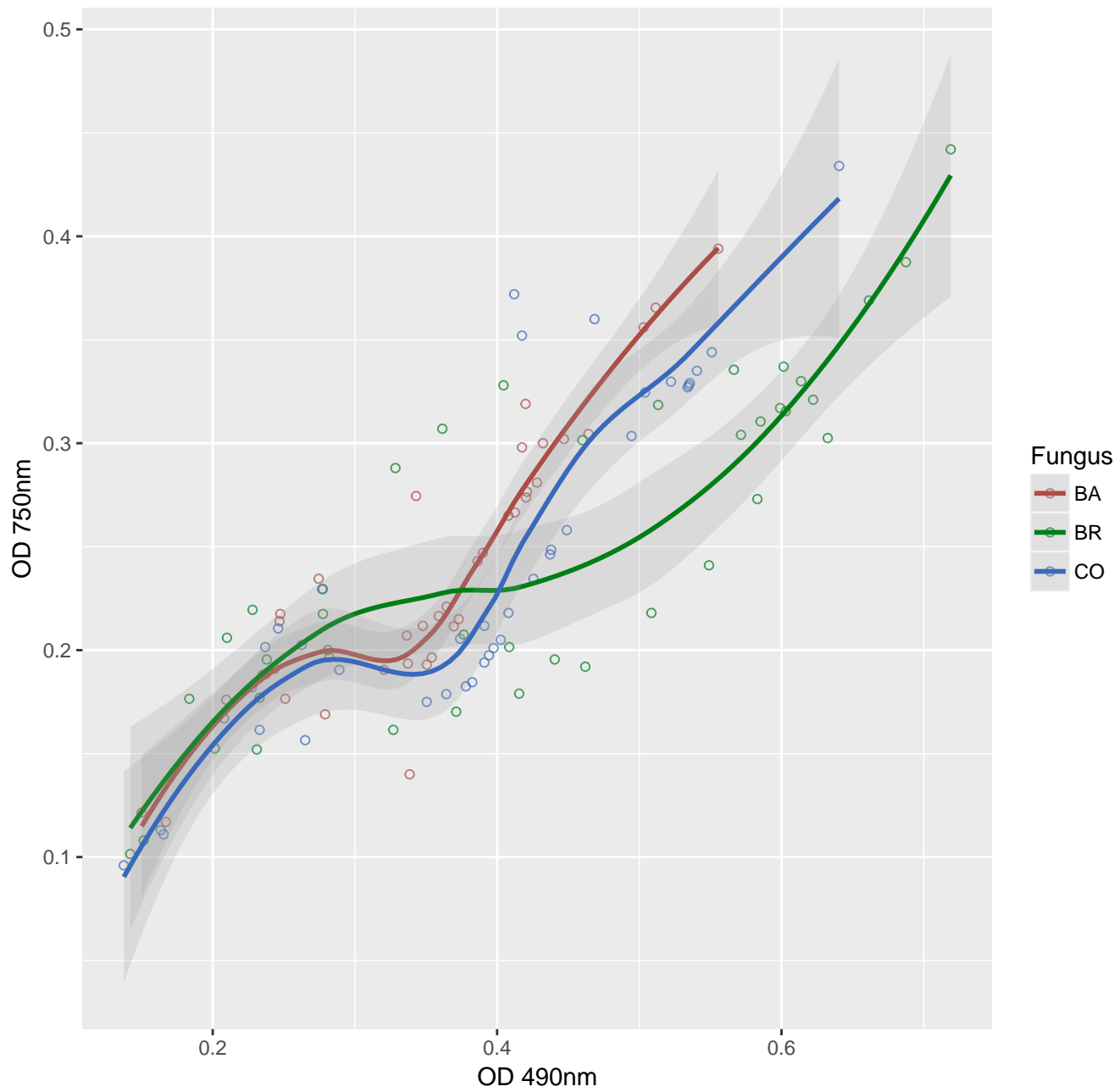
Adonitol



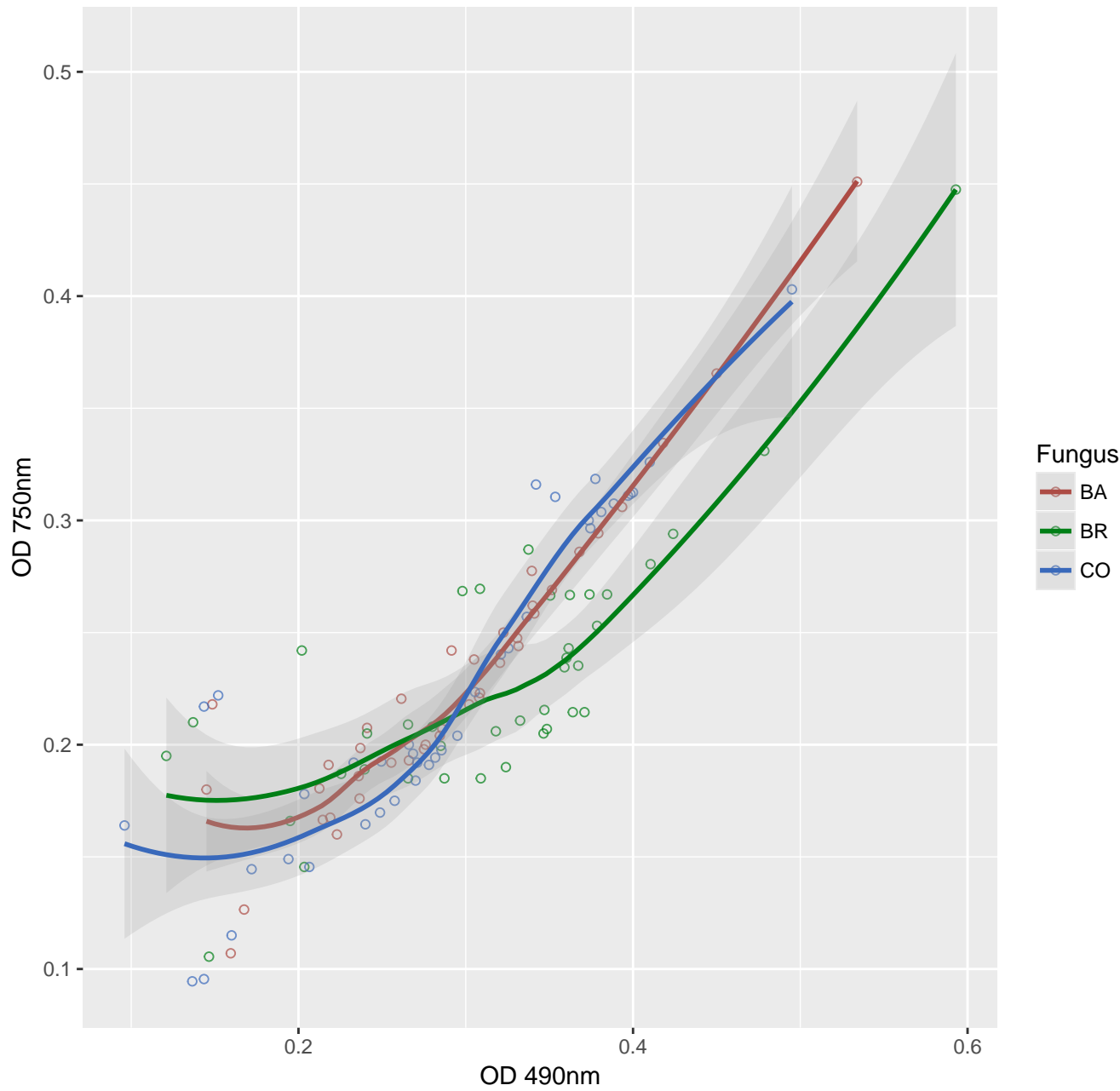
A.Keto.glutaricAcid



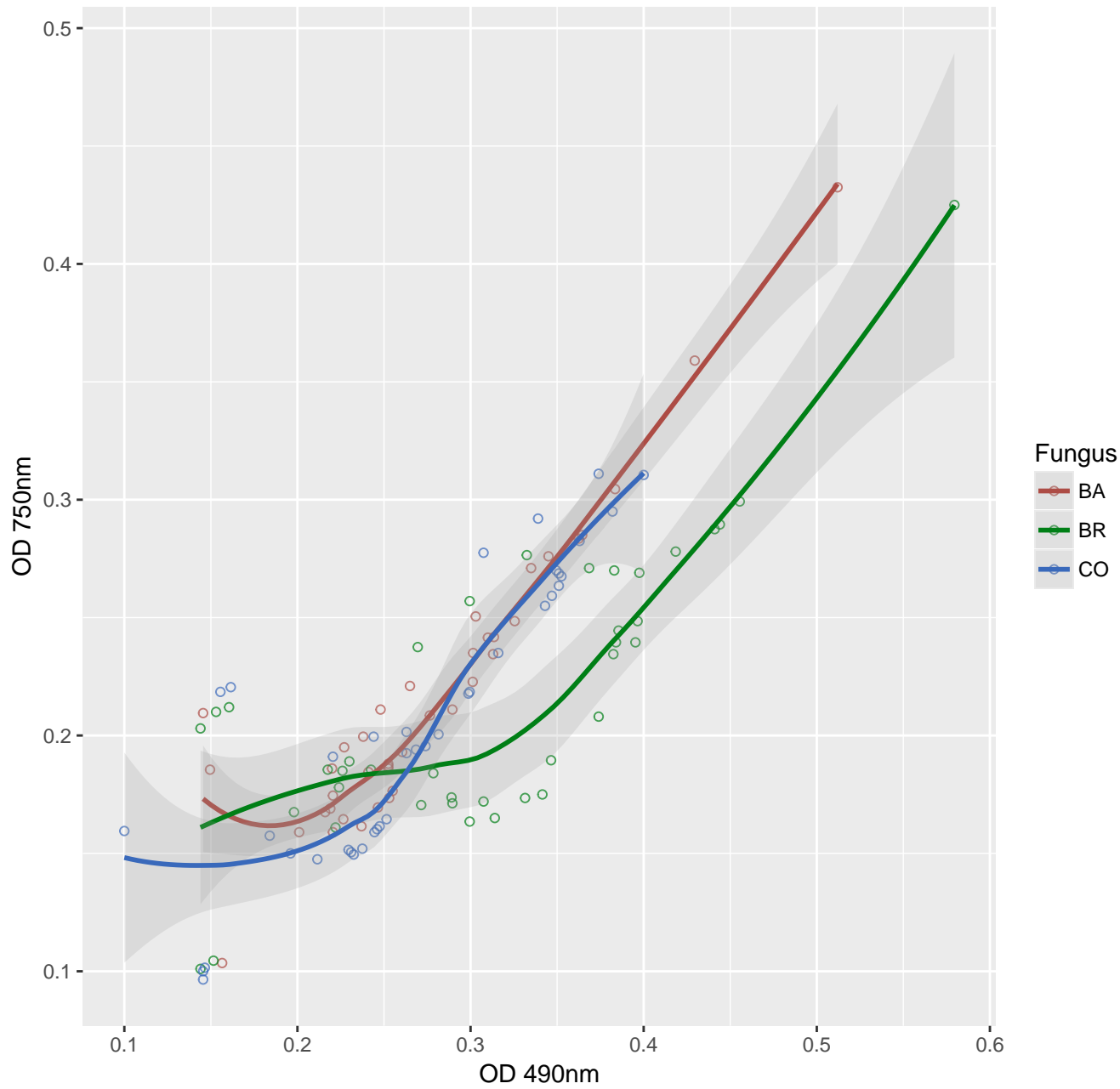
Alaninamide



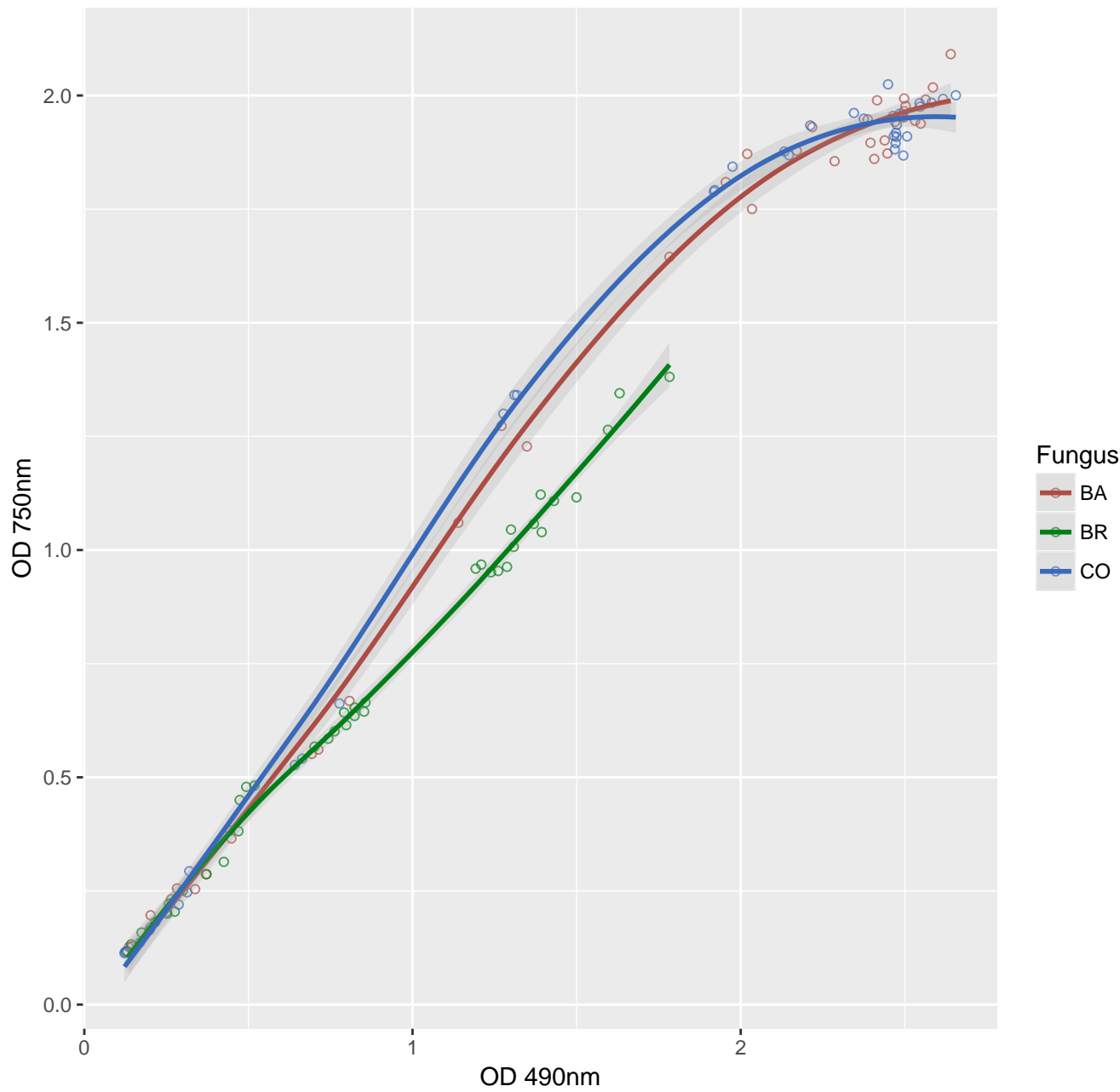
a.Methyl.D.Galactoside



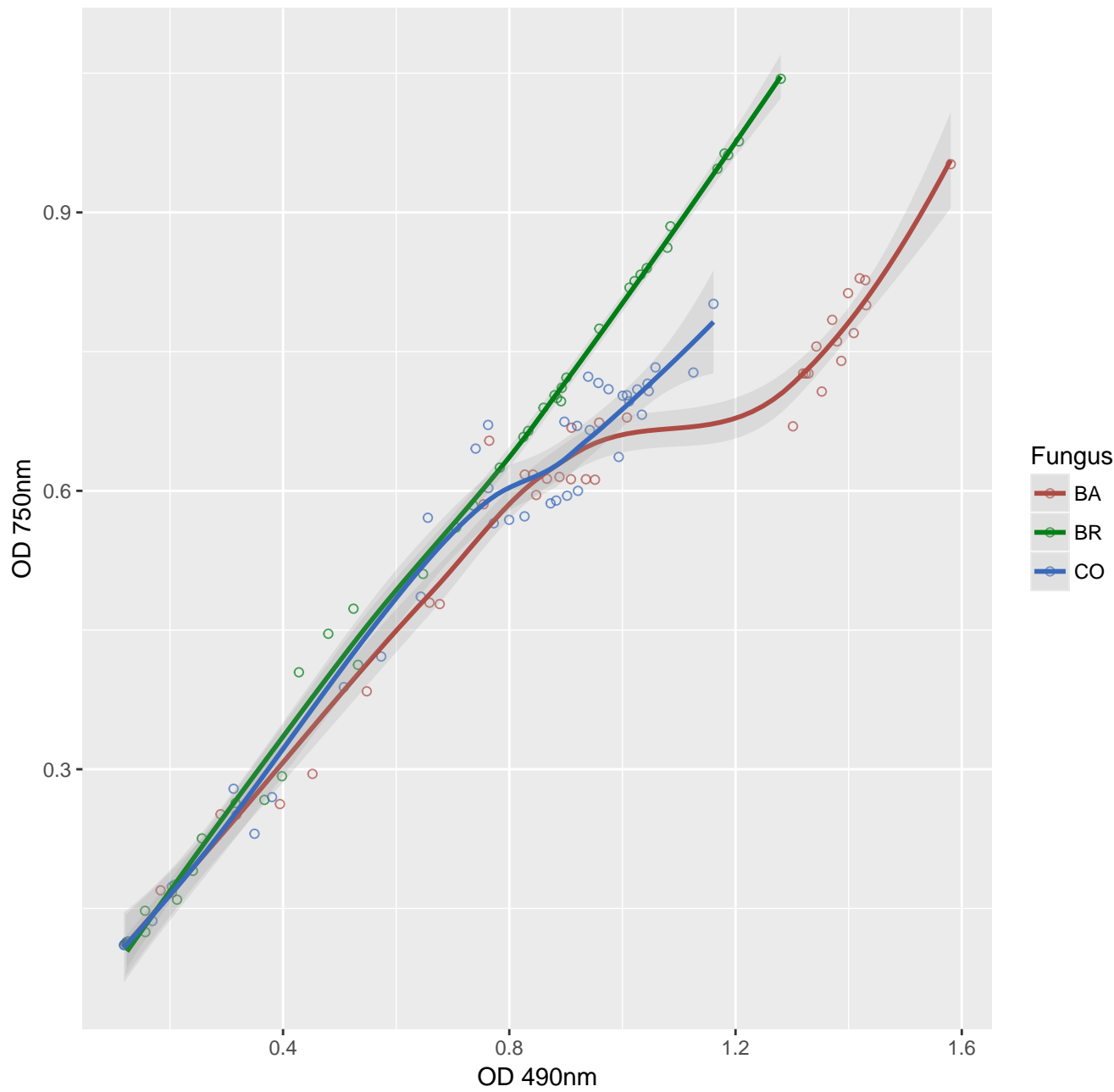
a.Methyl.D.Glucoside



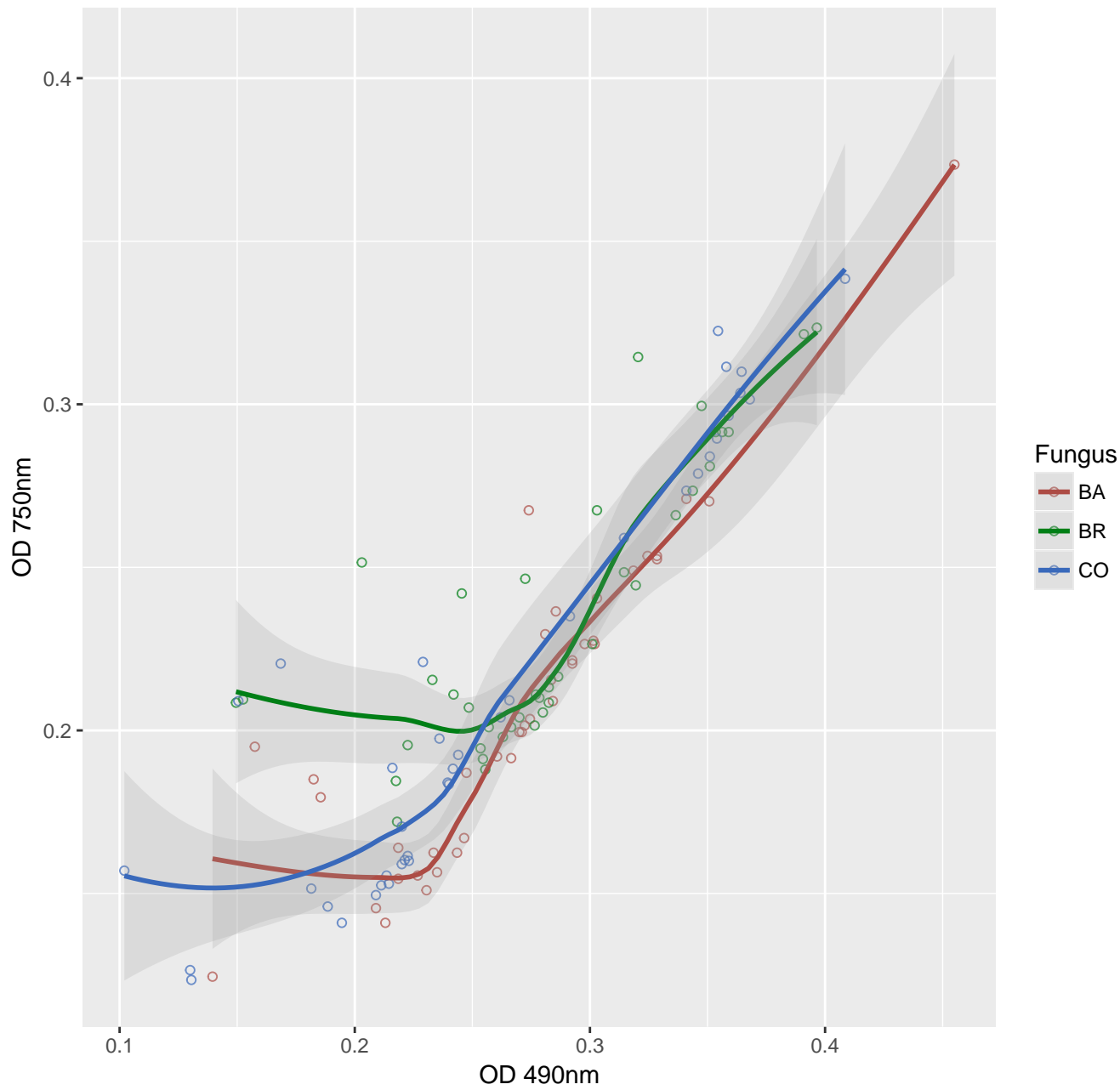
Amygdalin



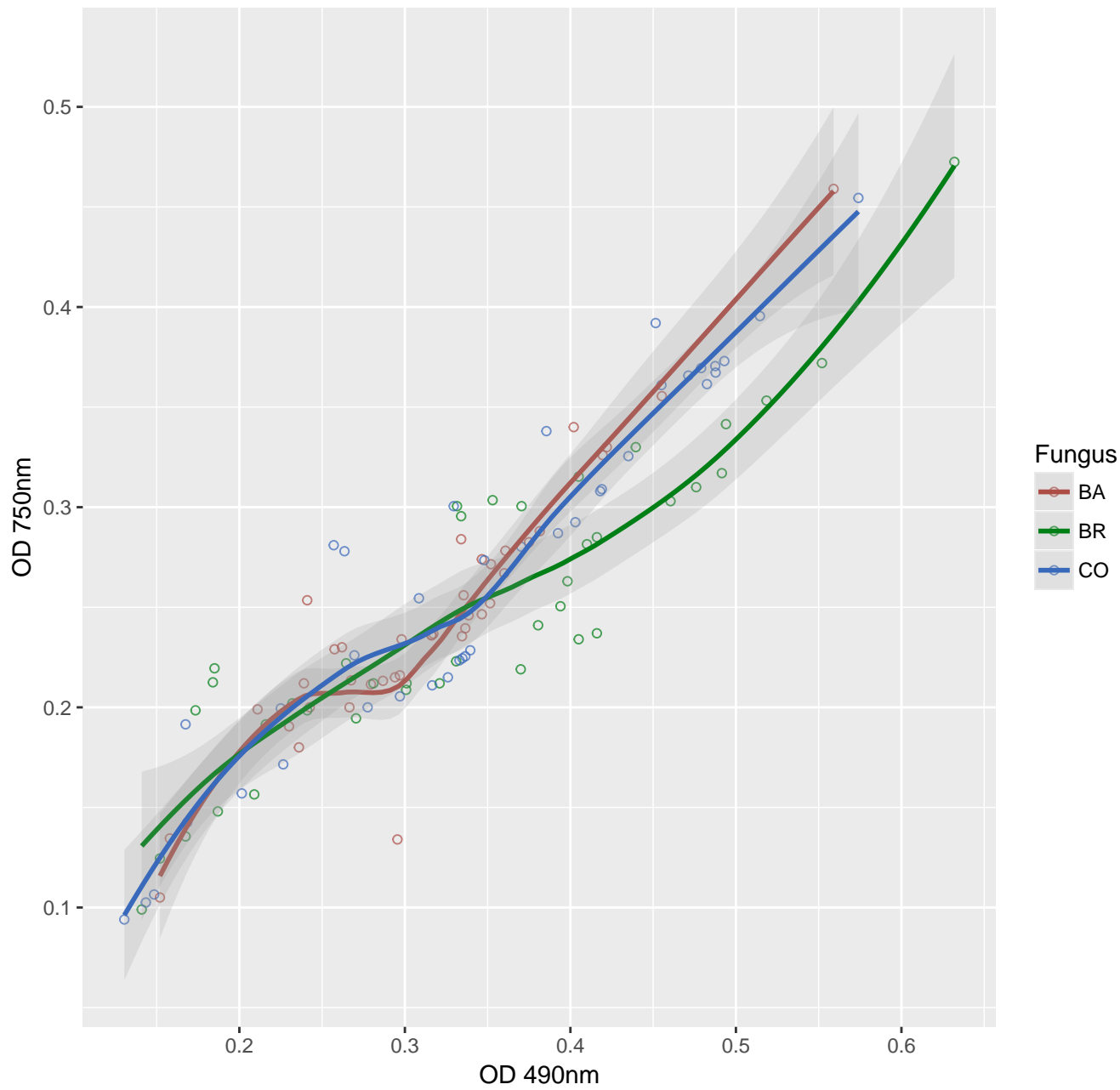
Arbutin



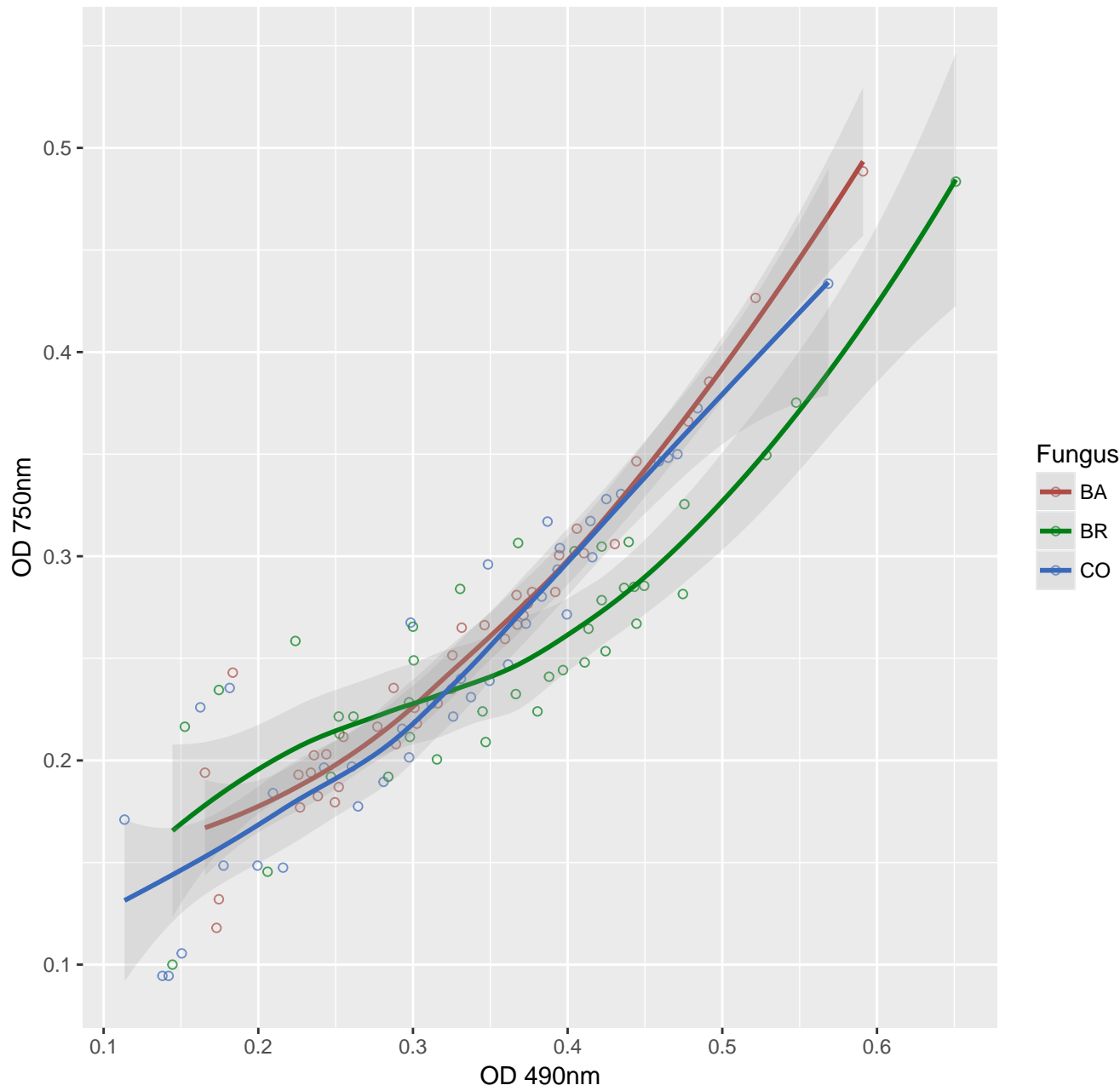
b. Cyclodextrin



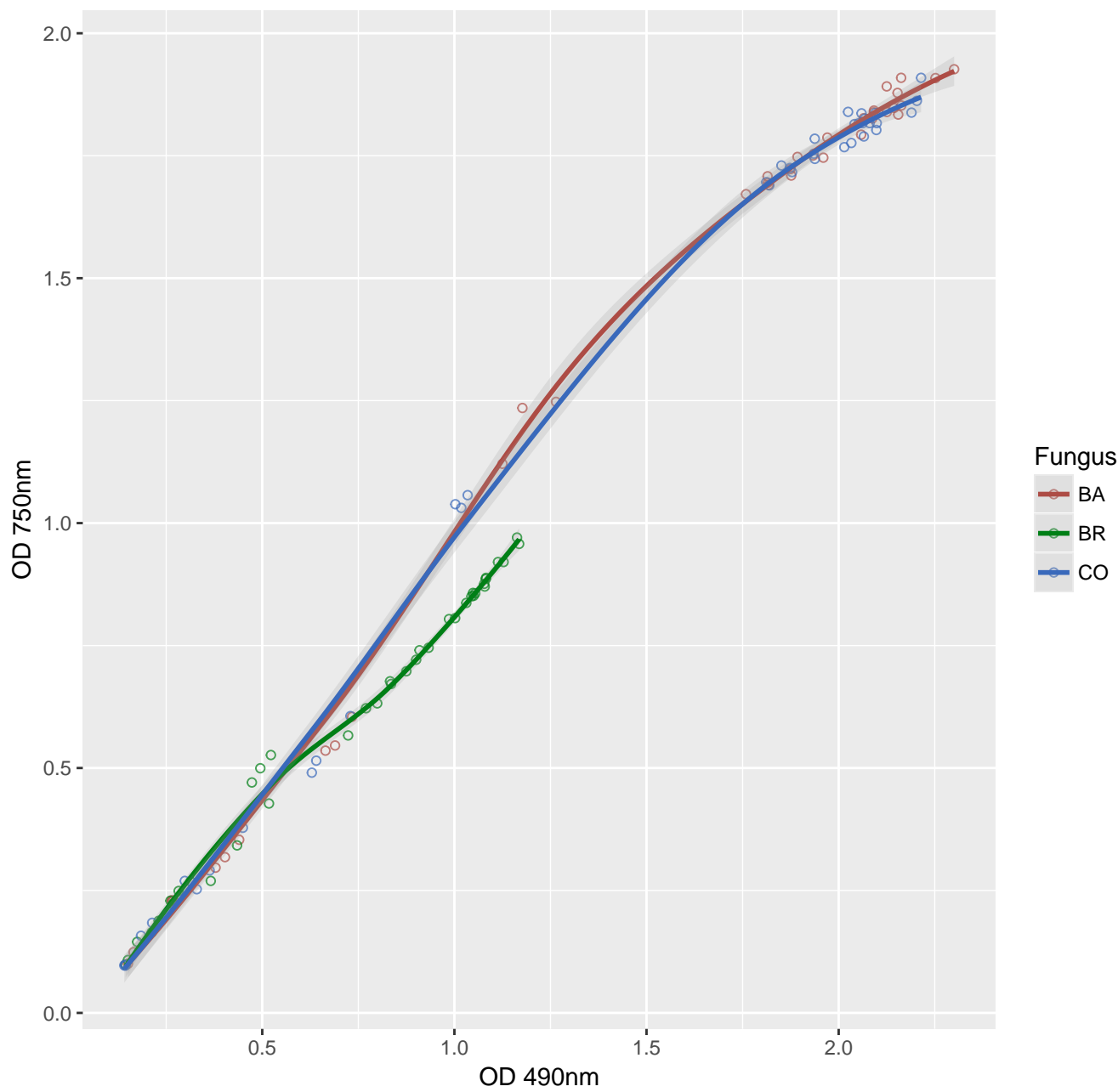
B. Hydroxy. butyric Acid



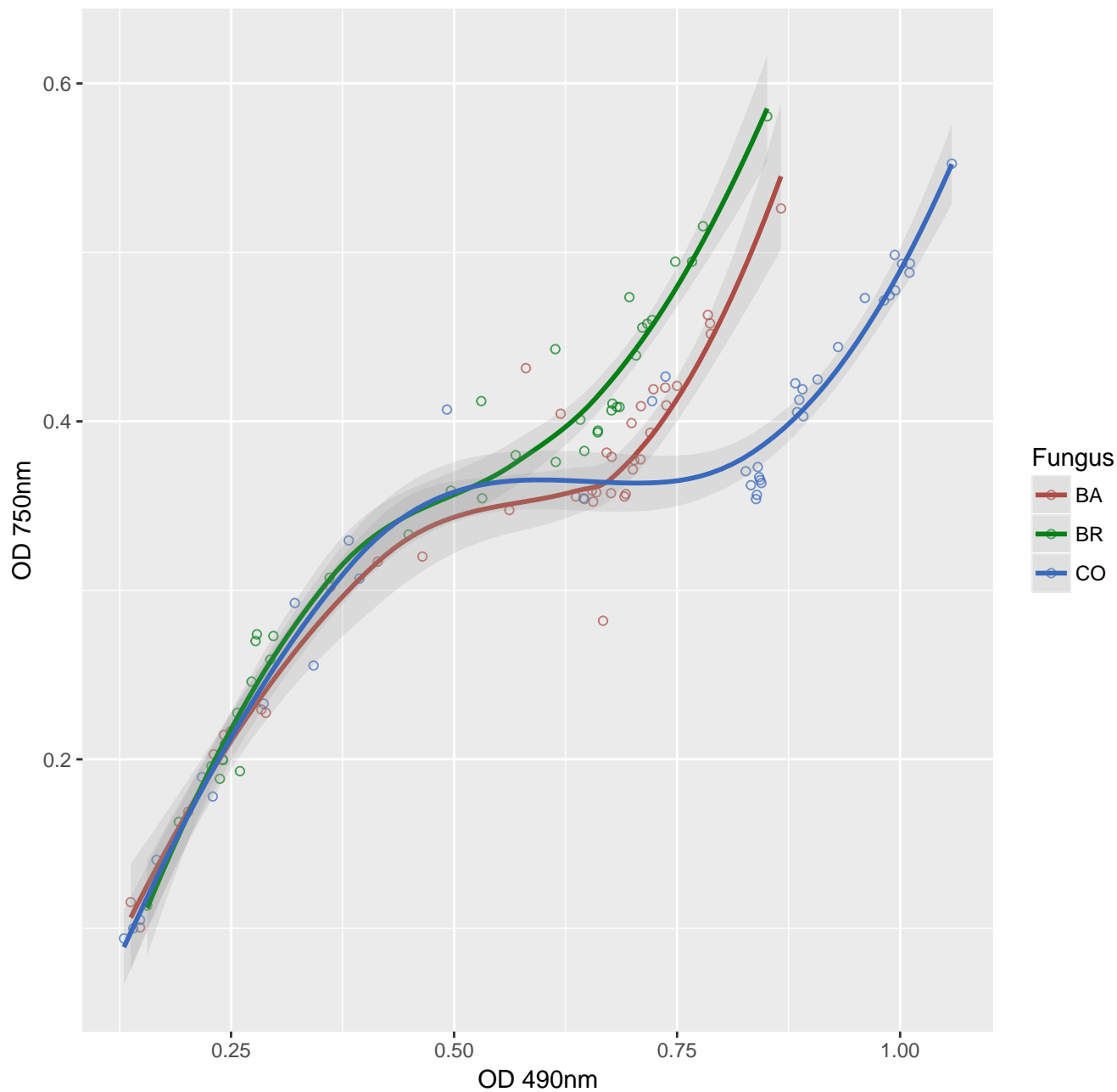
b.Methyl.D.Galactoside



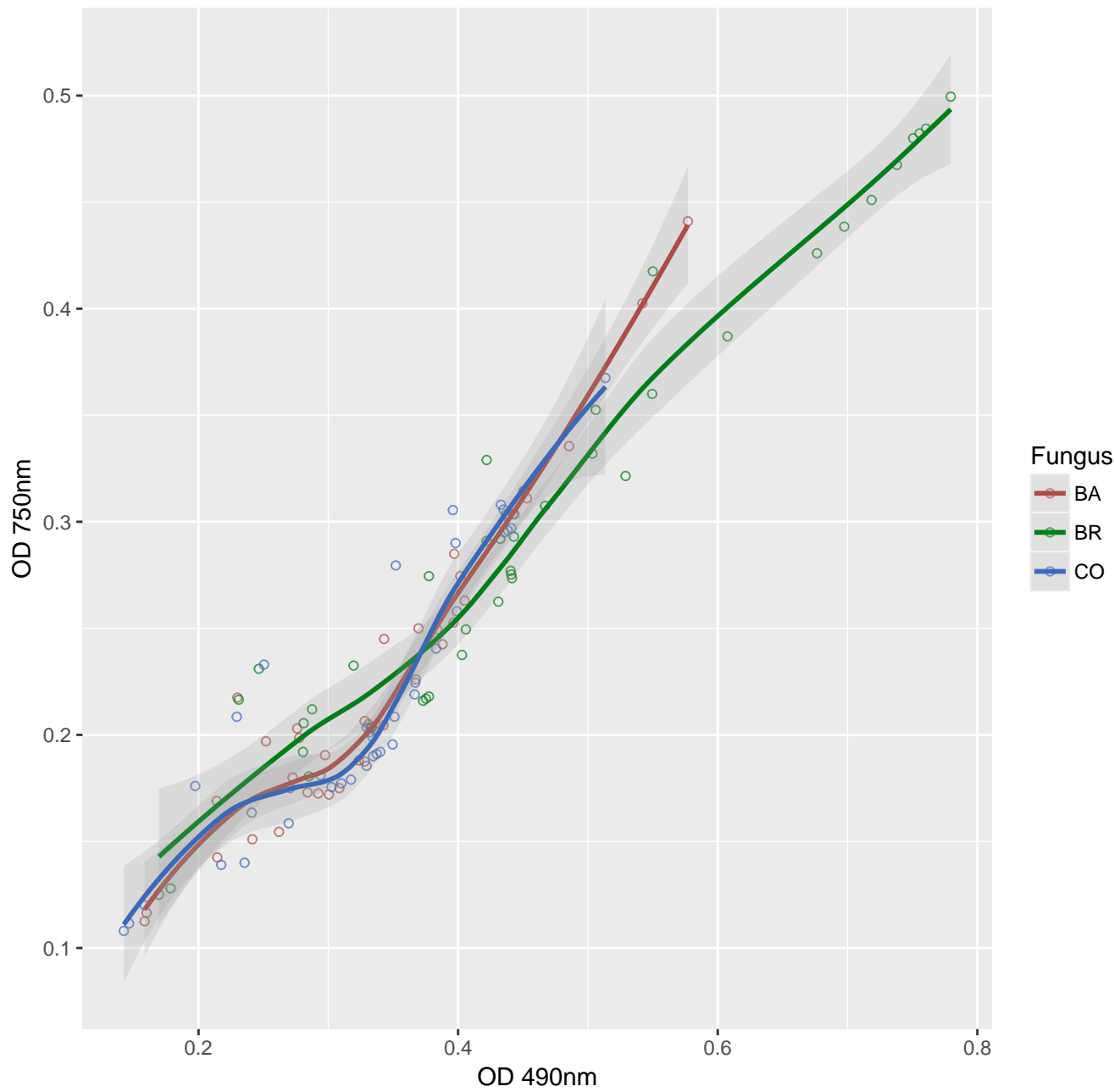
b.Methyl.D.Glucoside



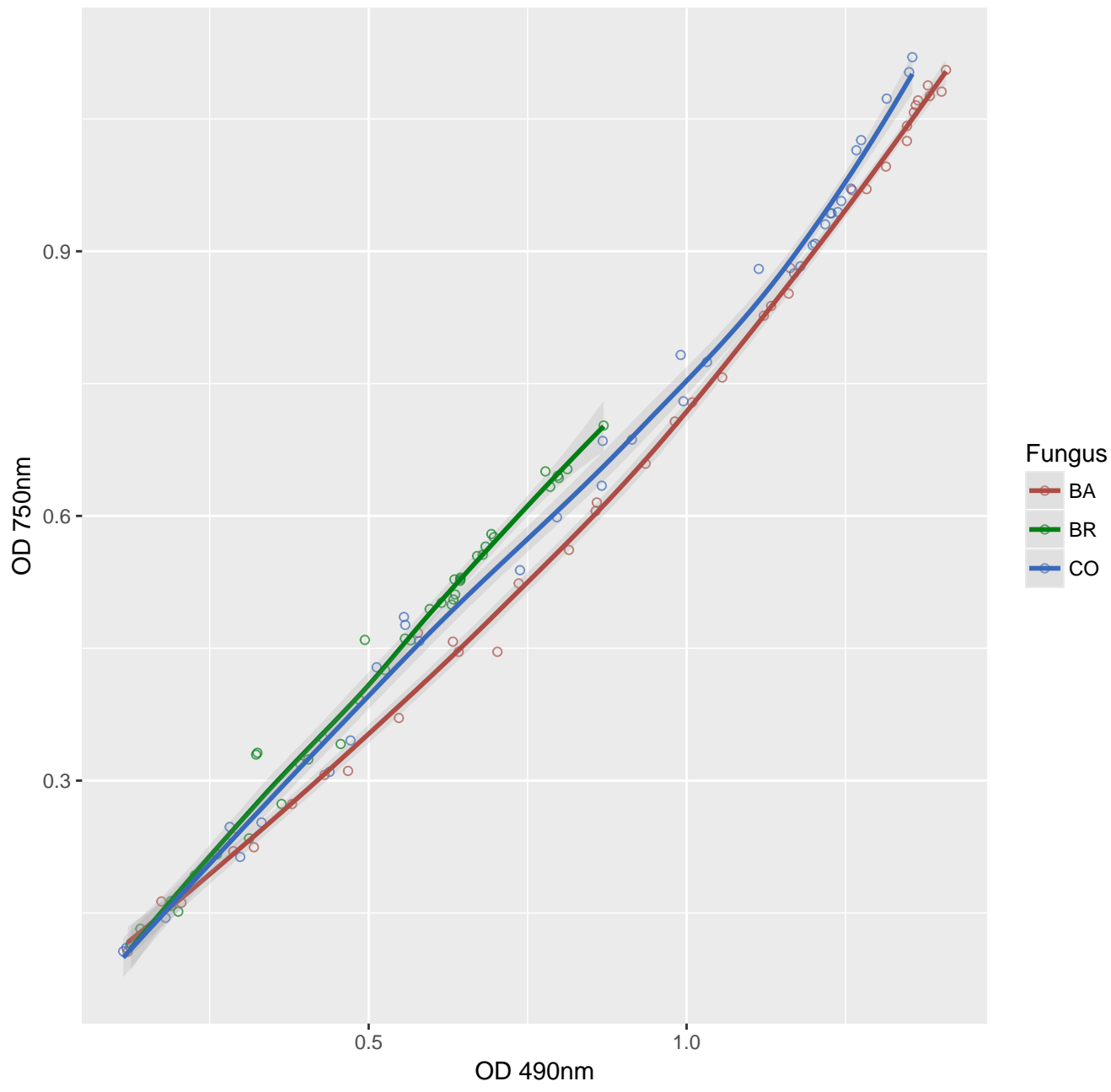
BromosuccinicAcid



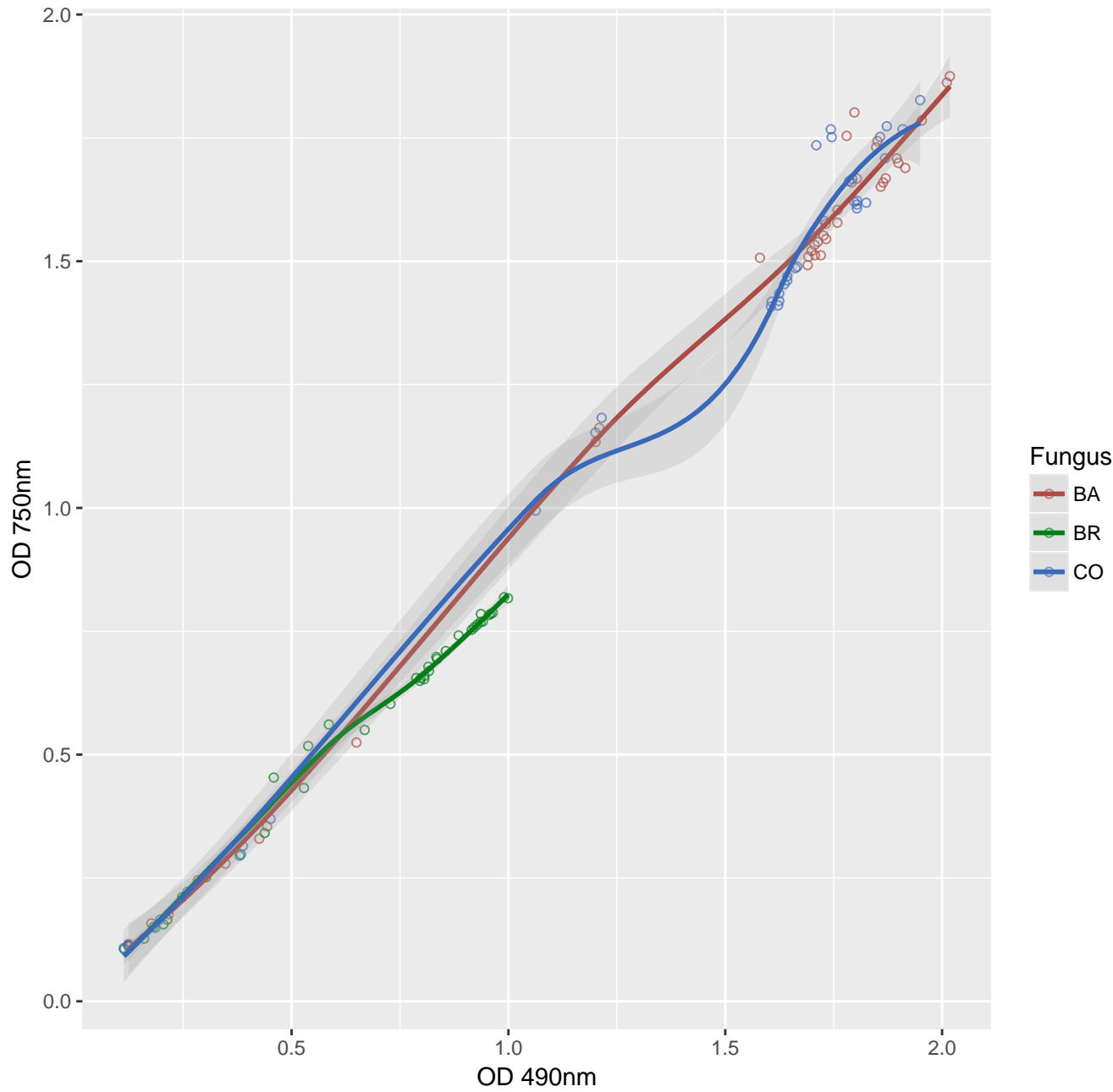
D.Arabinose



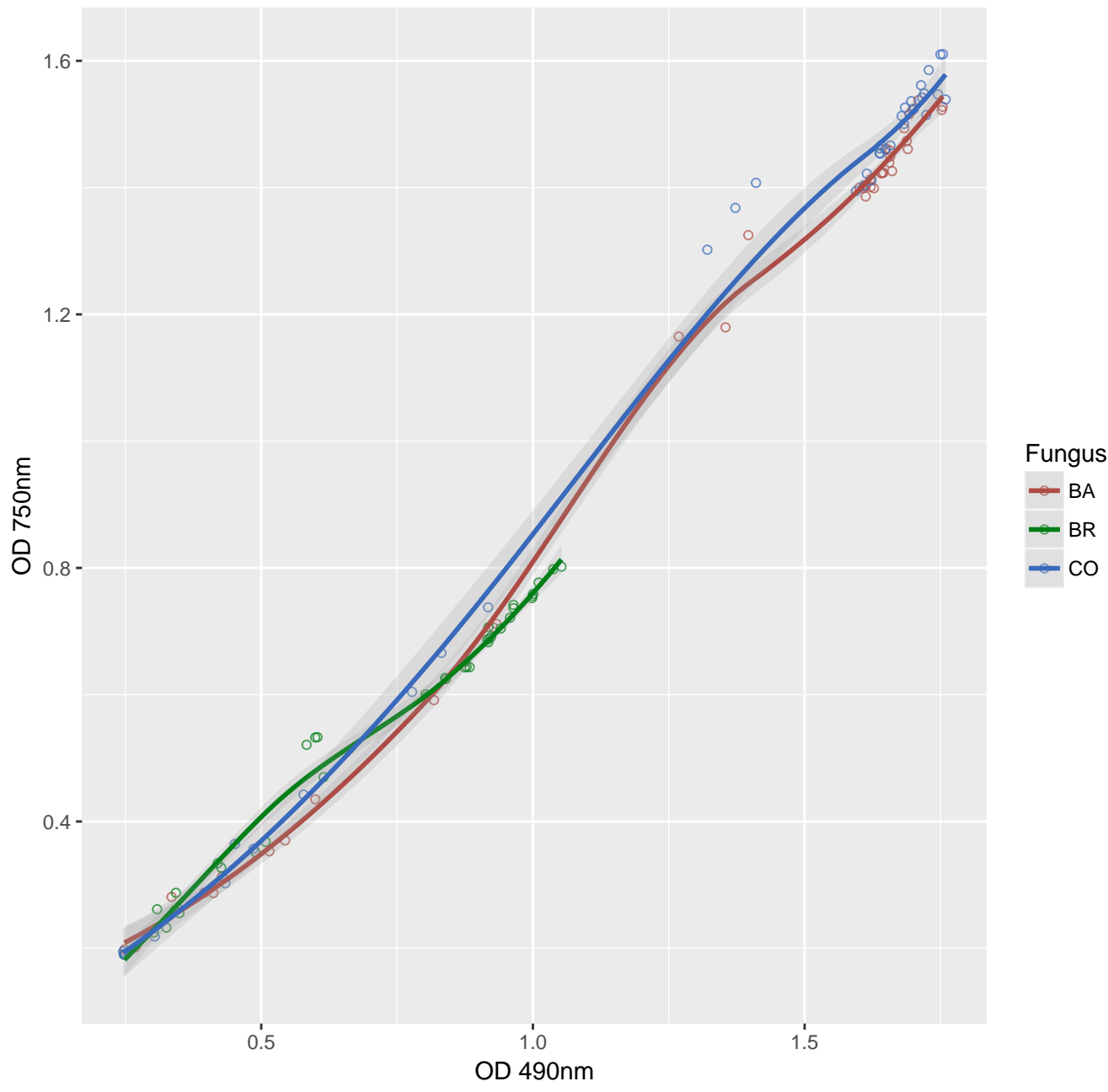
D.Arabitol



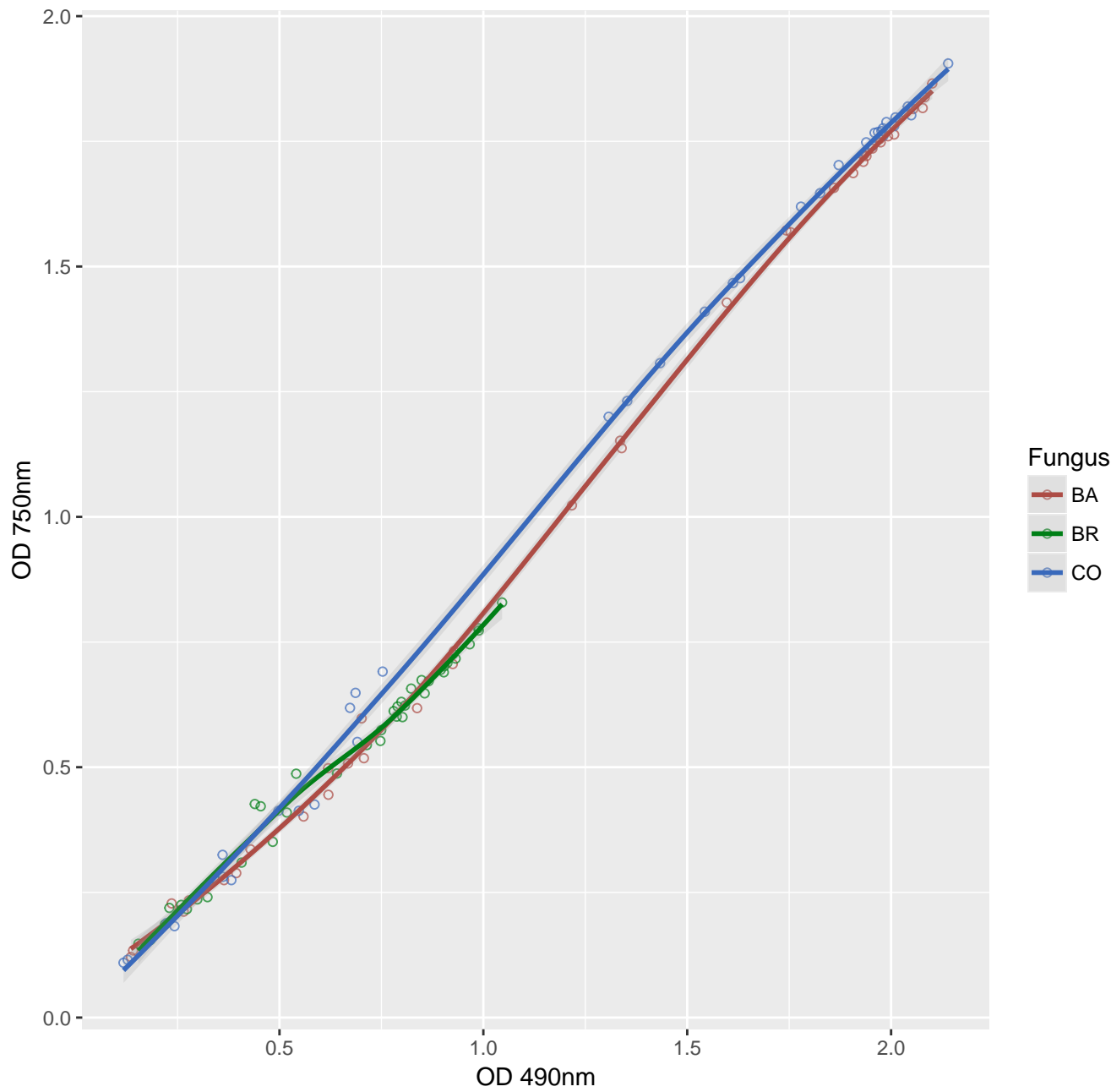
D.Cellobiose



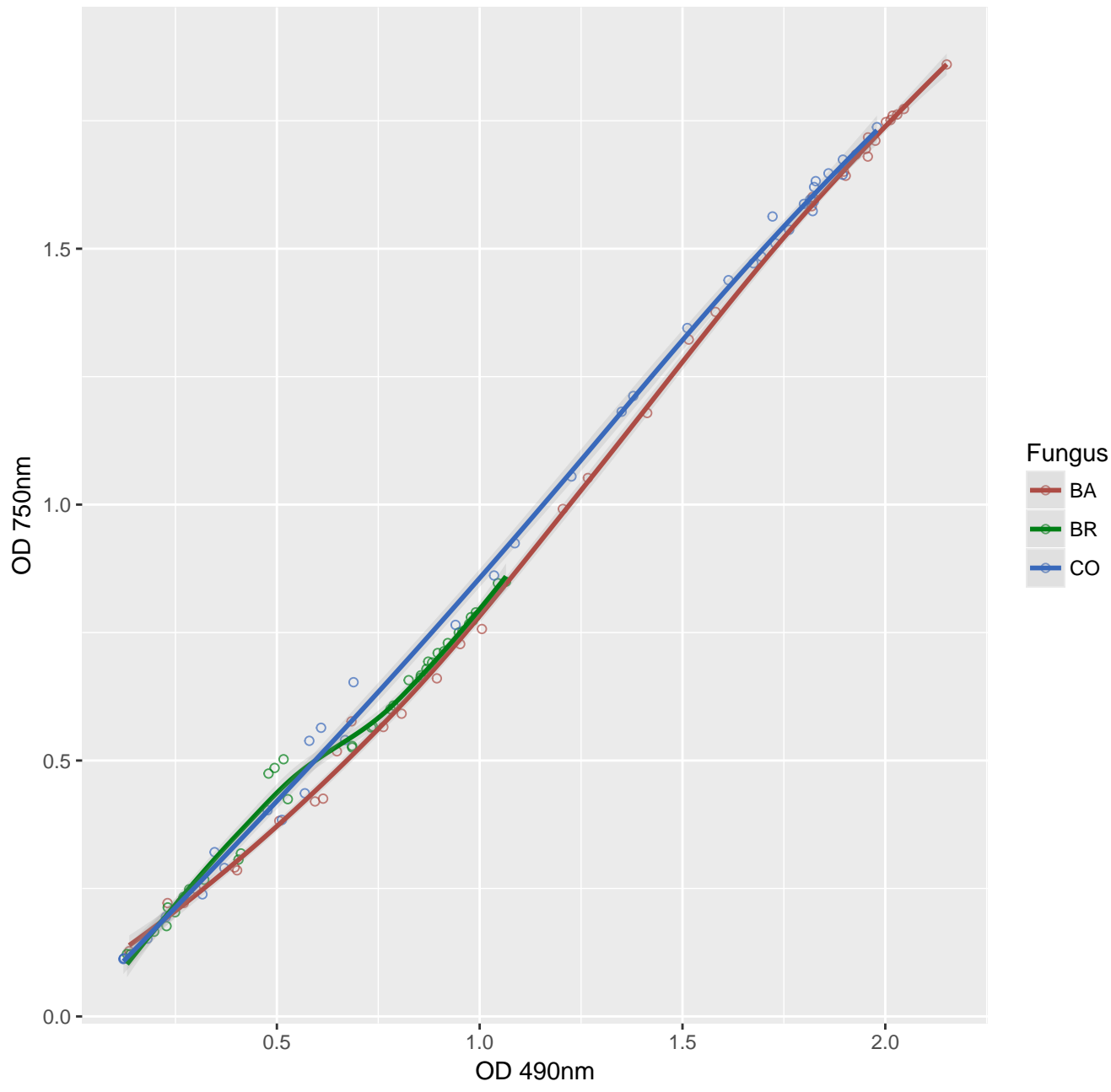
Dextrin



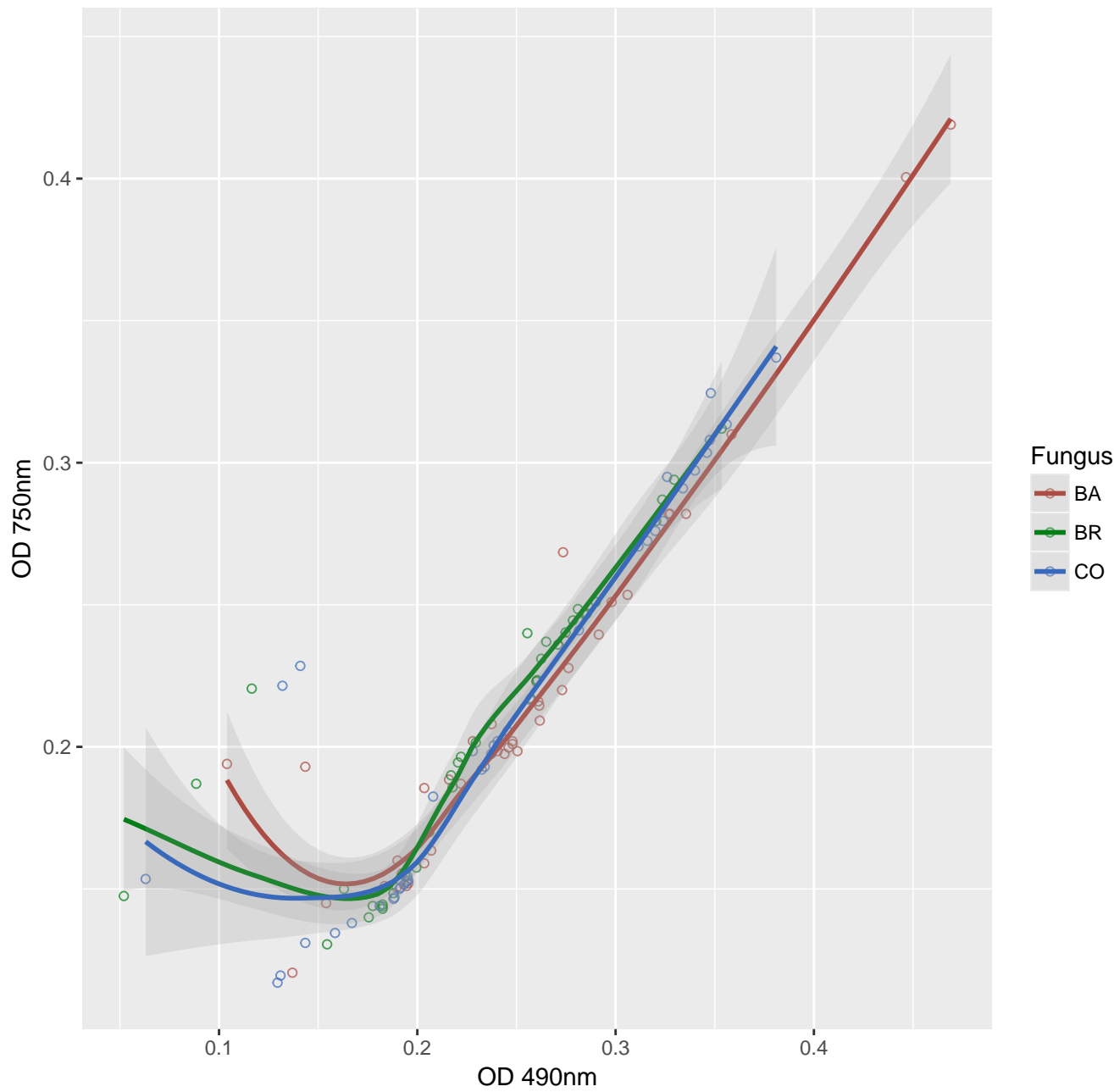
D.Fructose



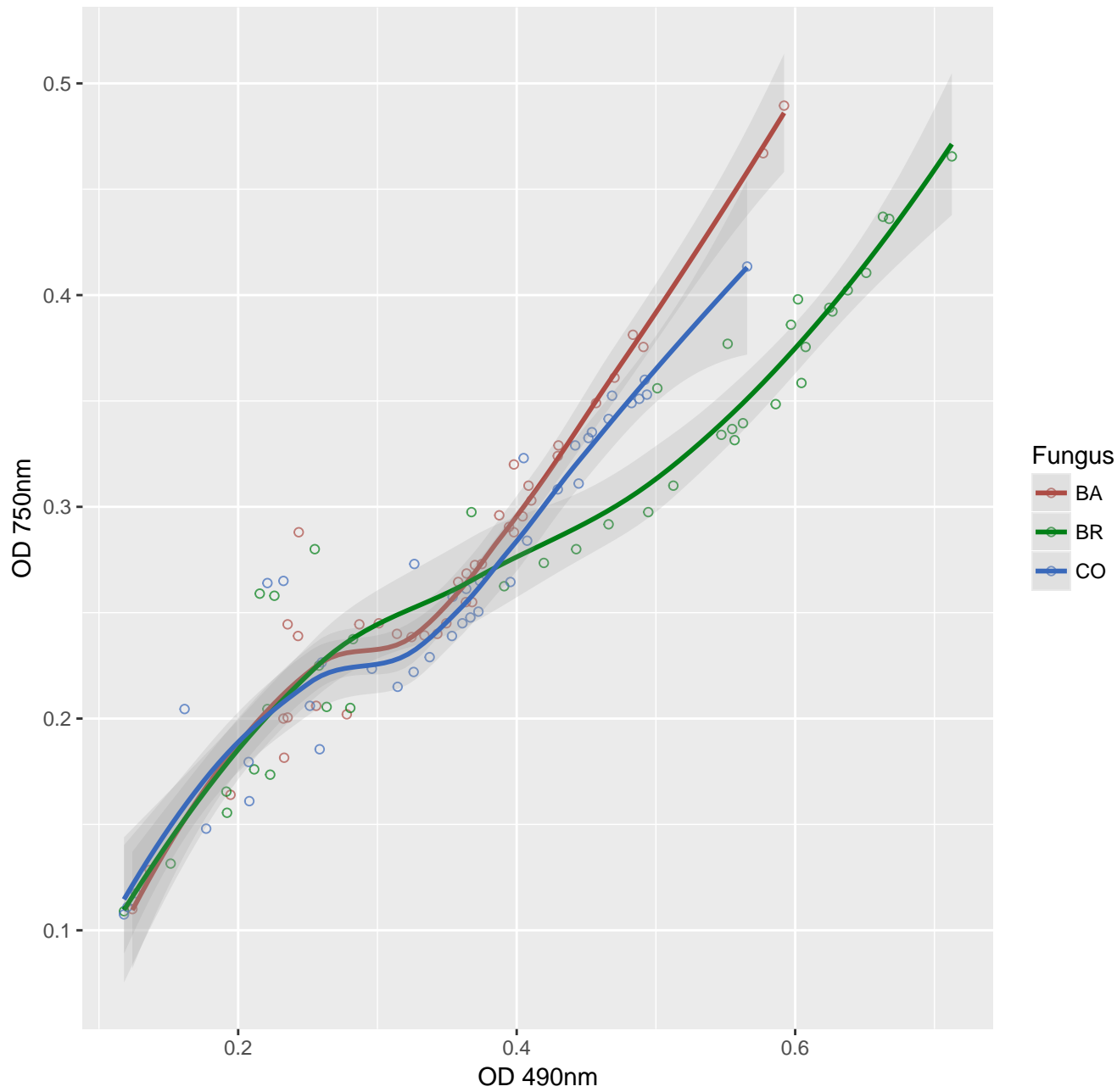
D.Galactose



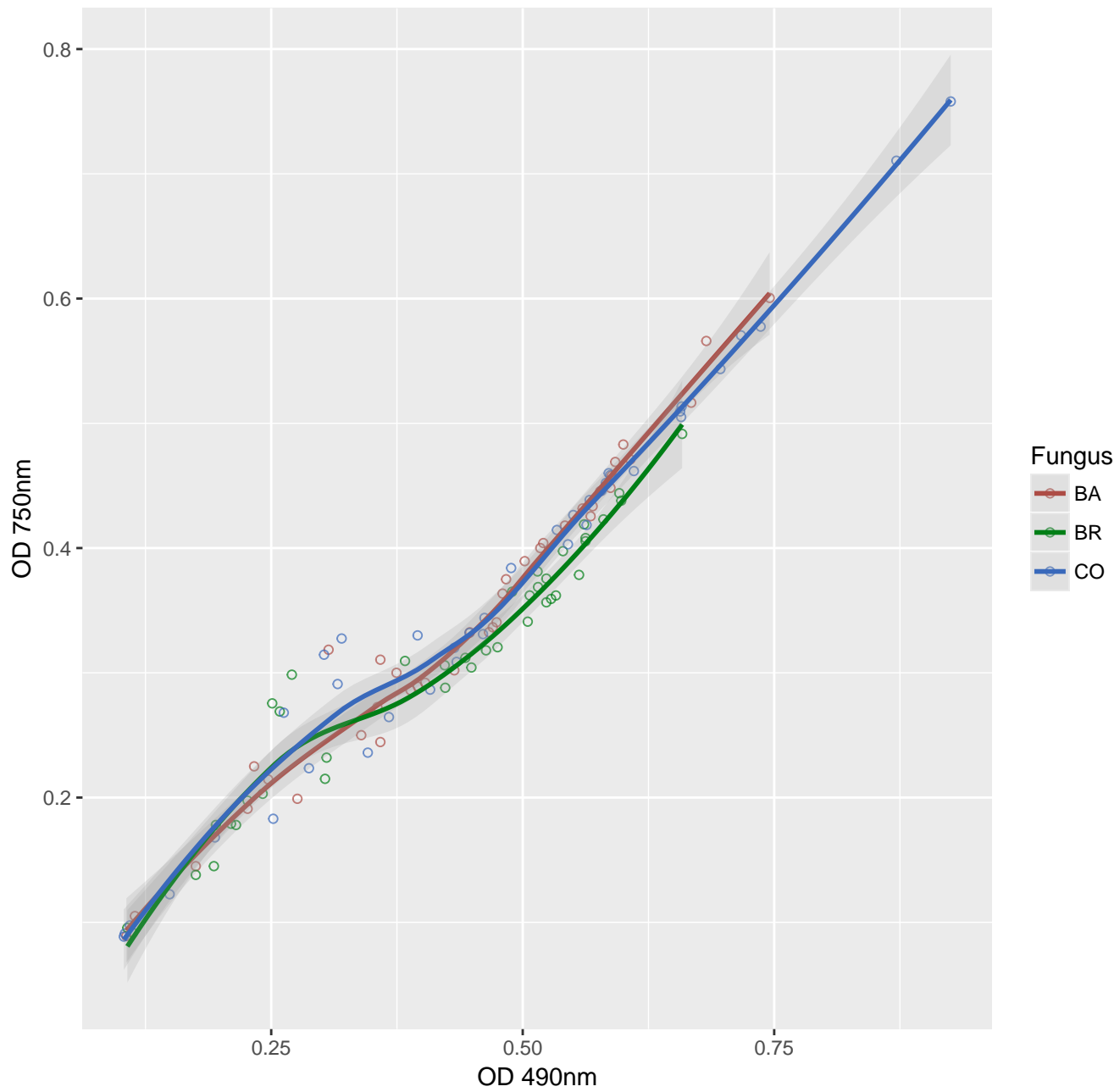
D.GalacturonicAcid



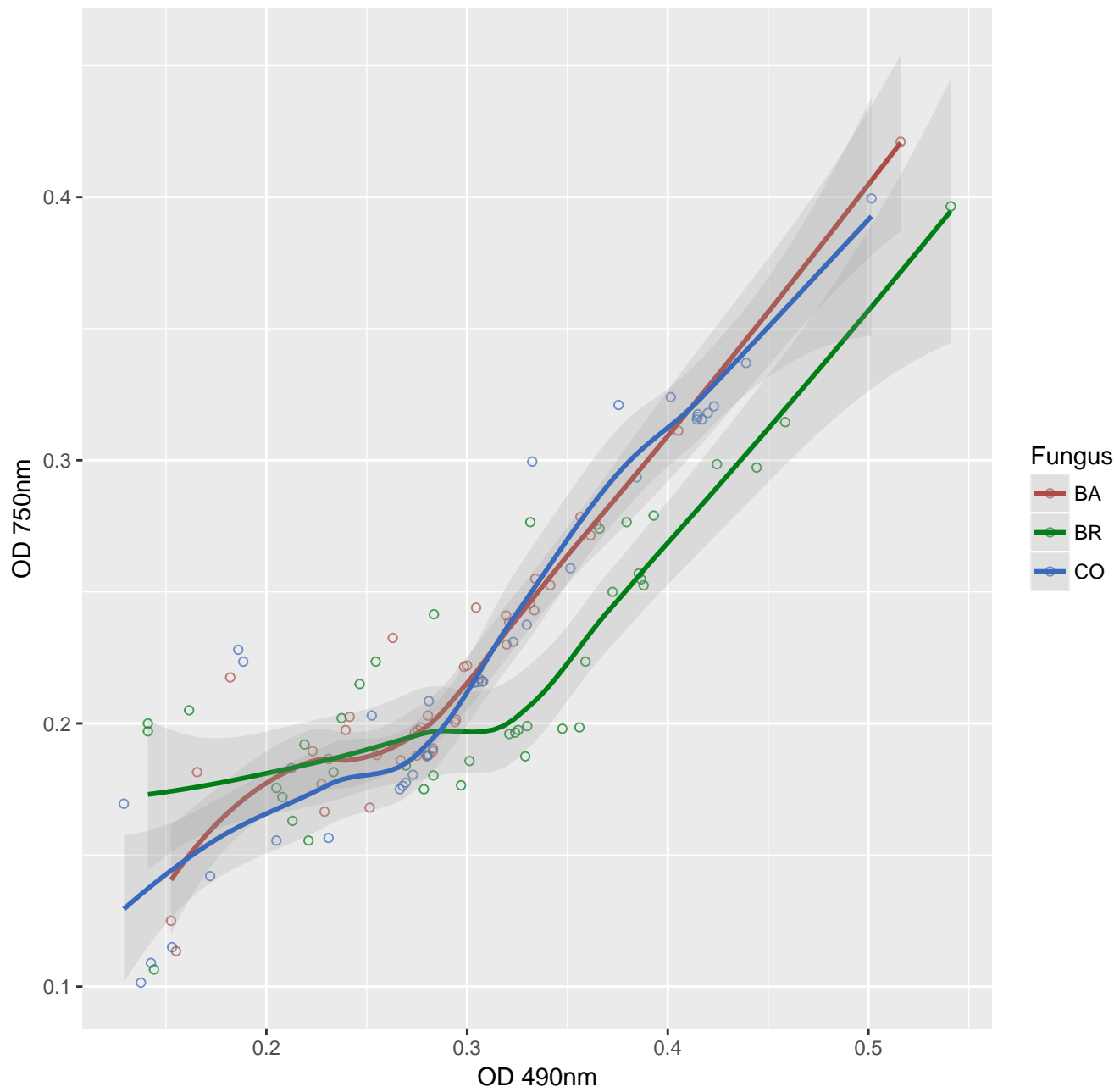
D.GluconicAcid



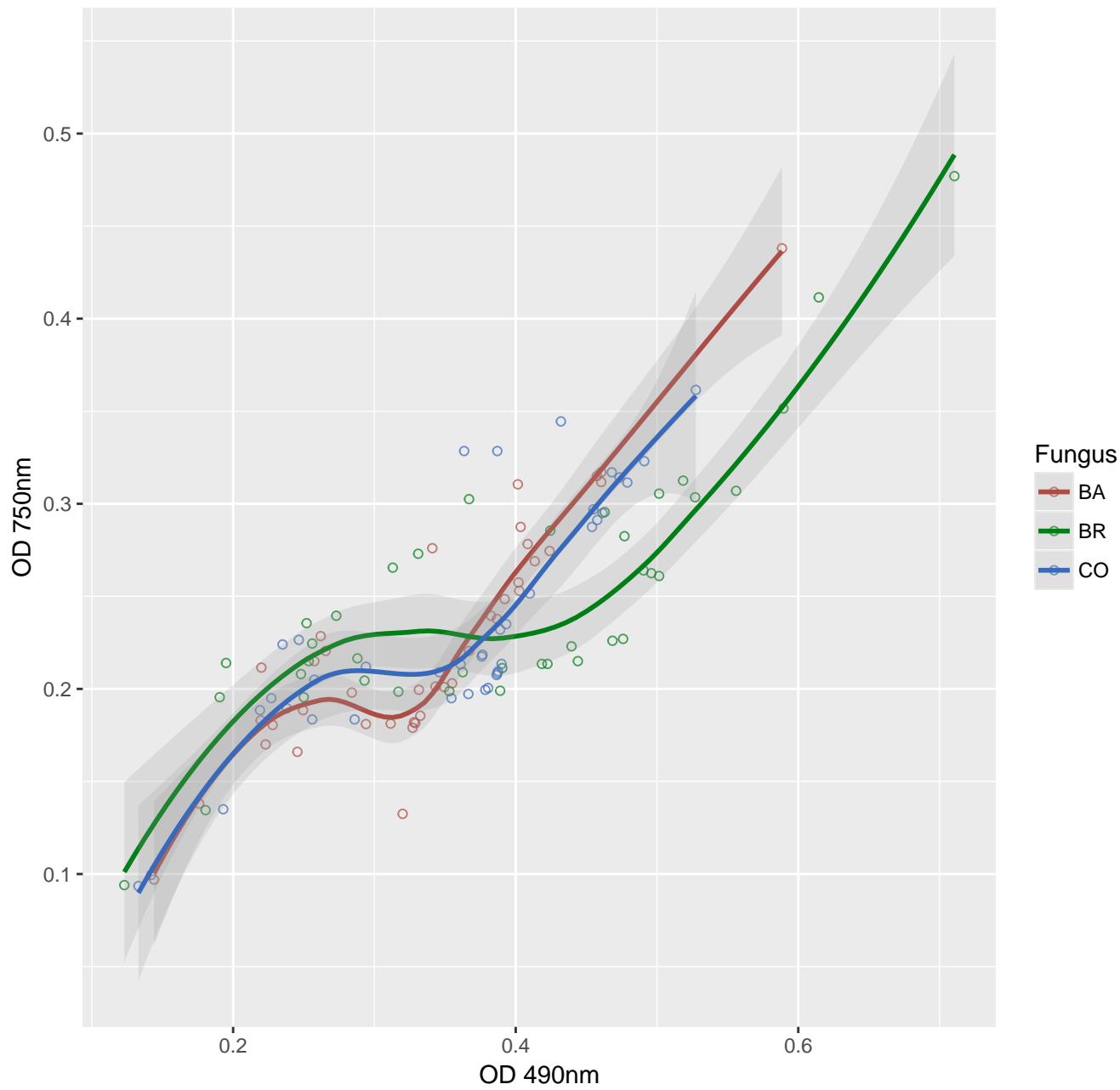
D.Glucosamine



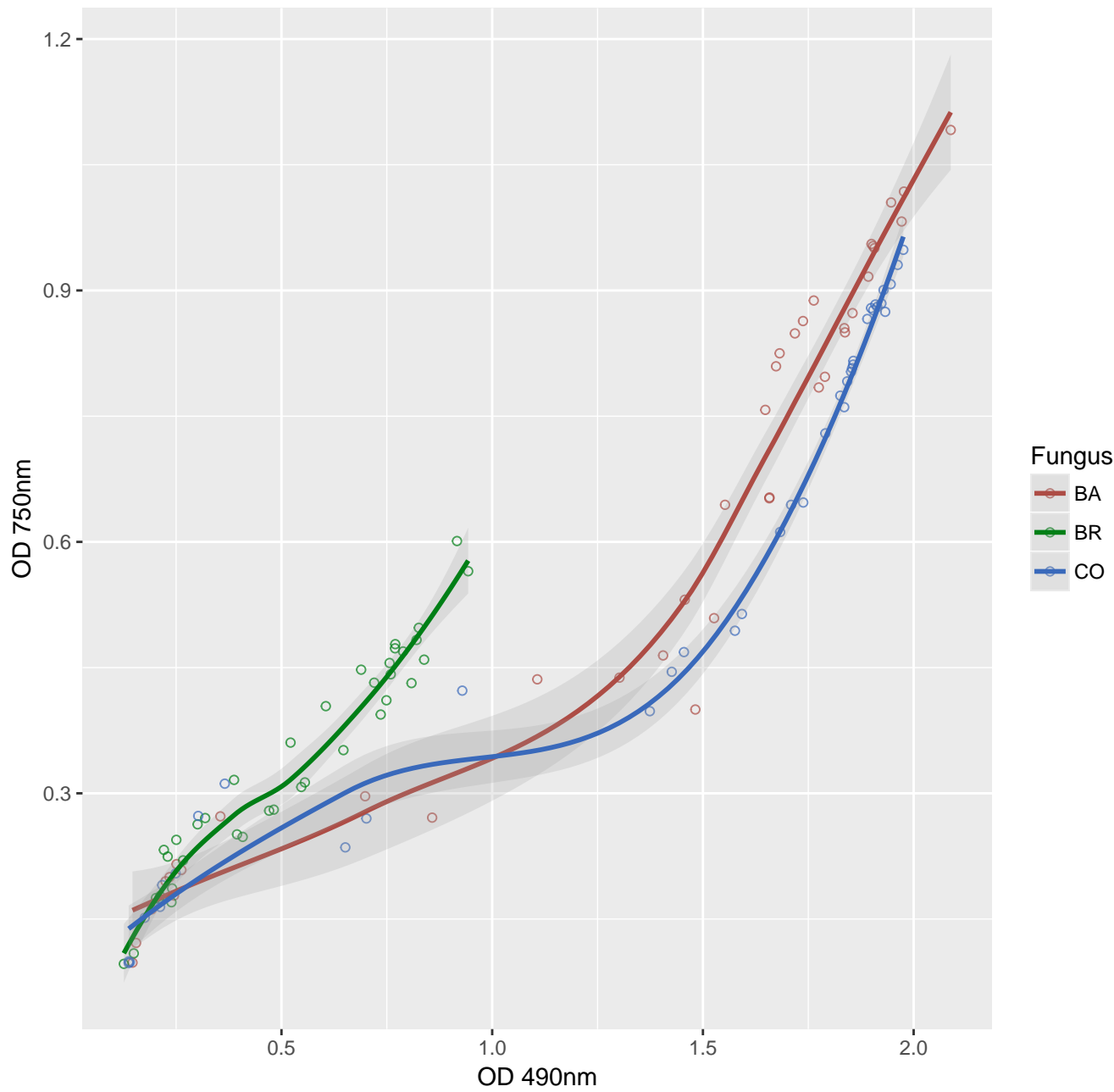
D.GlucuronicAcid



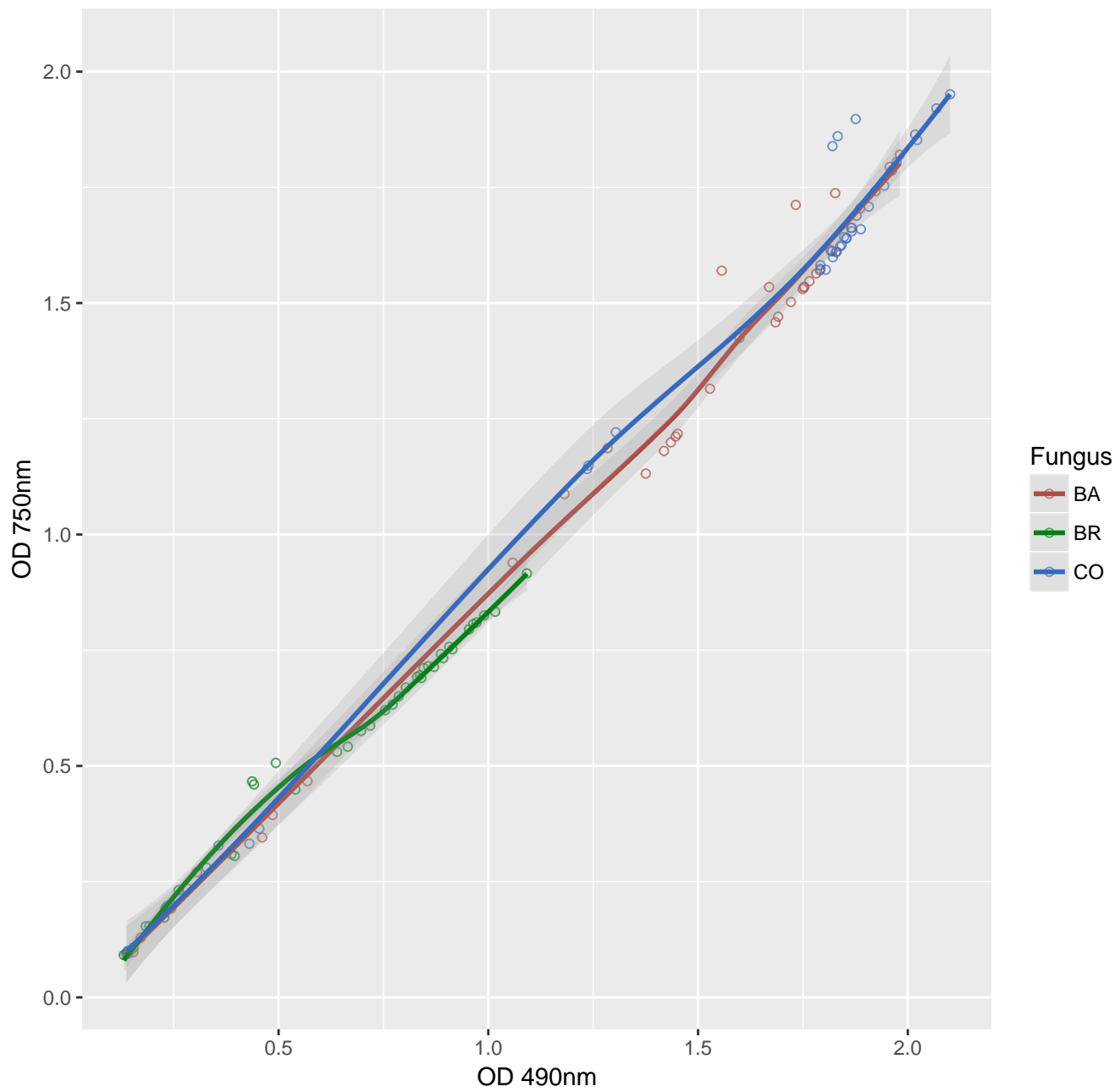
D.LacticAcidMethylEster



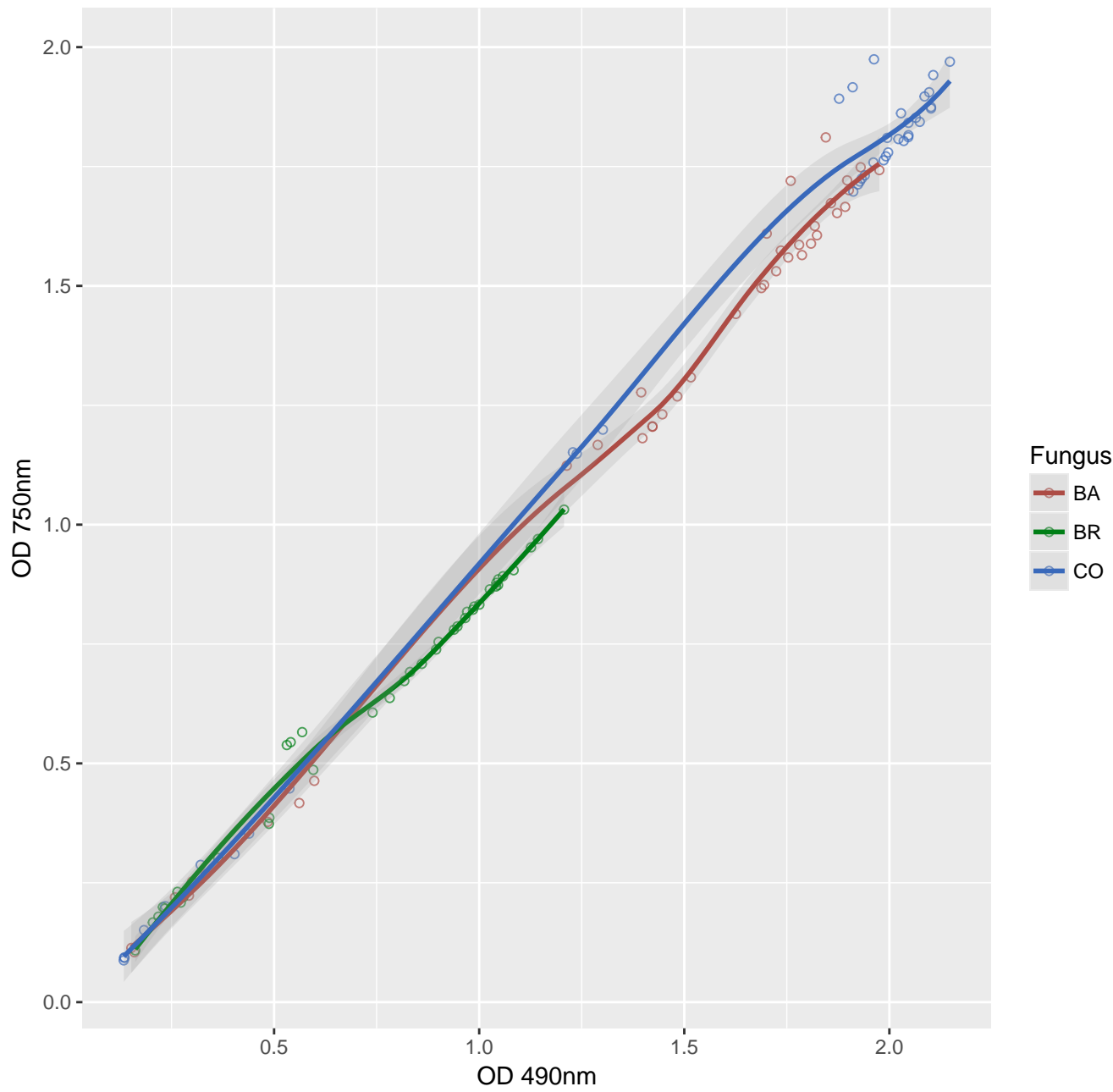
D.MalicAcid



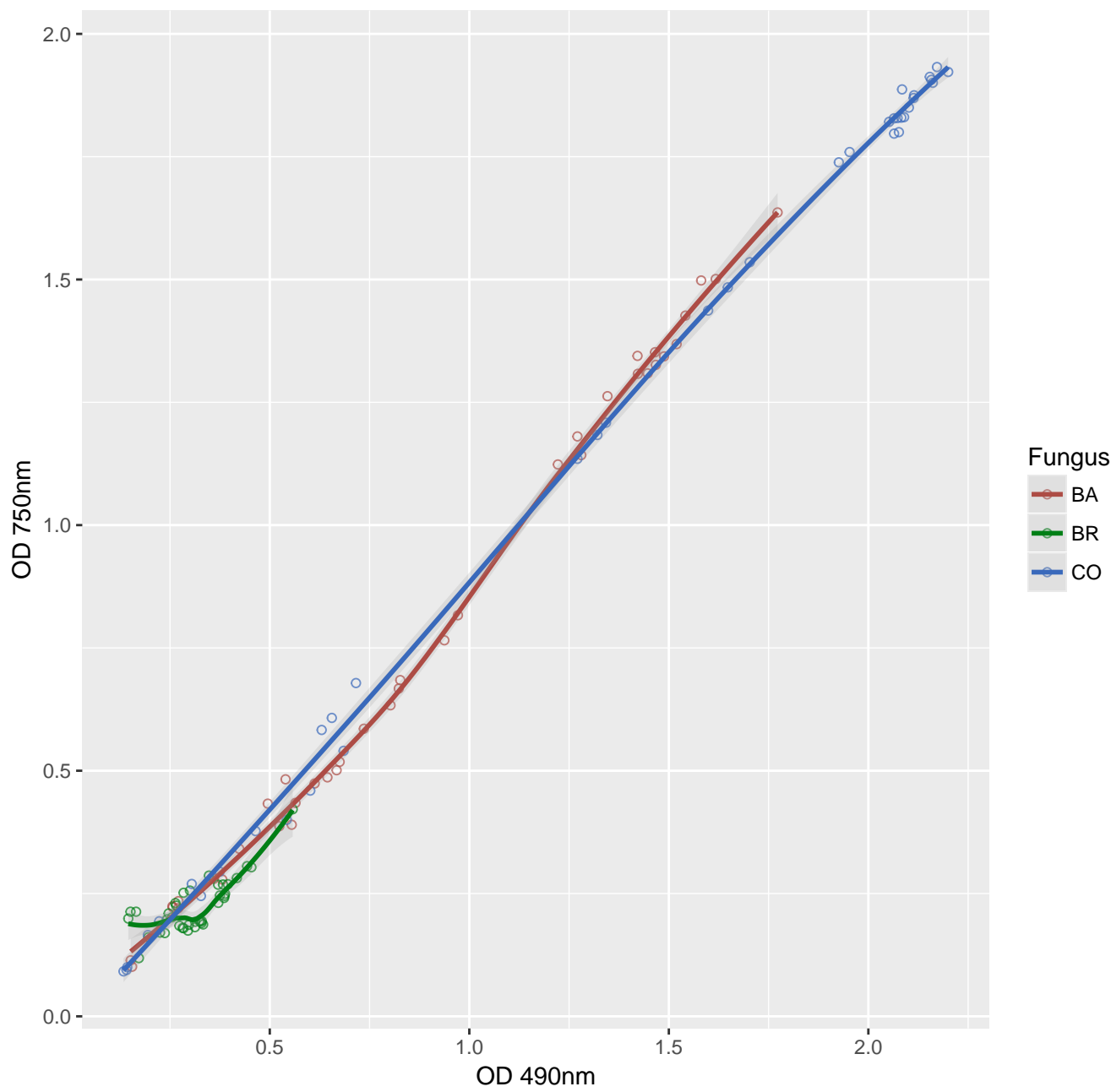
D.Mannitol



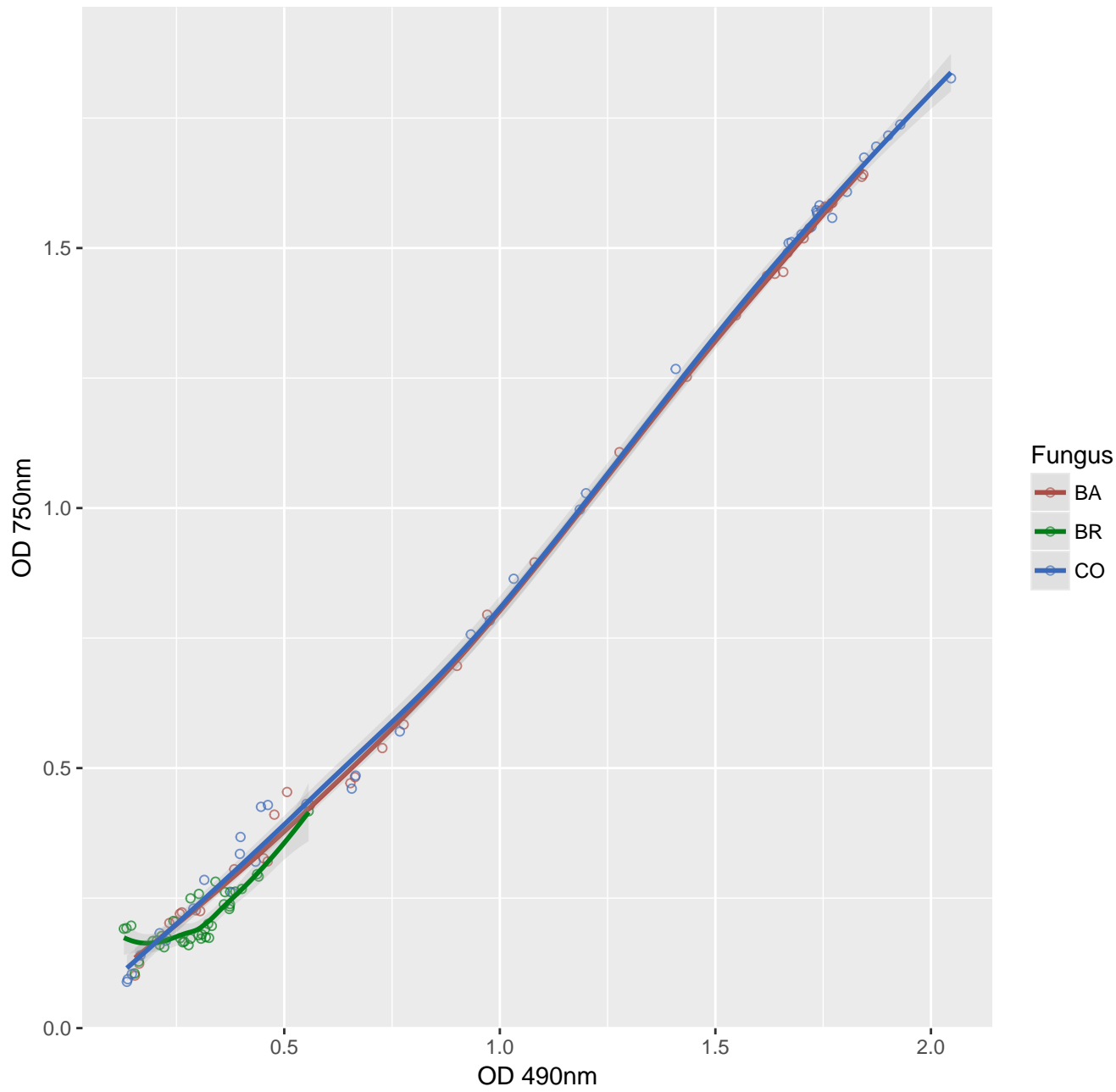
D.Mannose



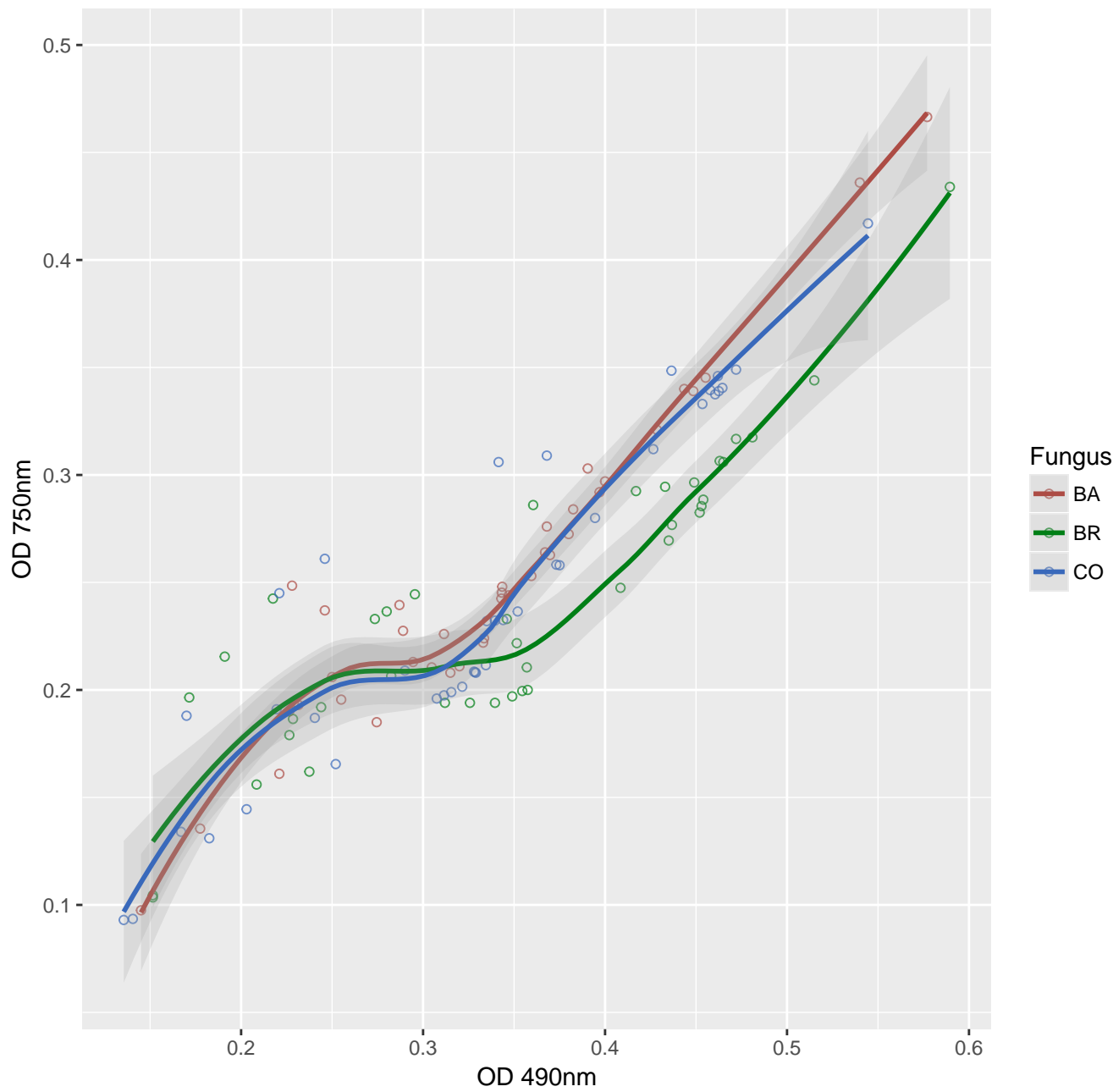
D.Melezitose



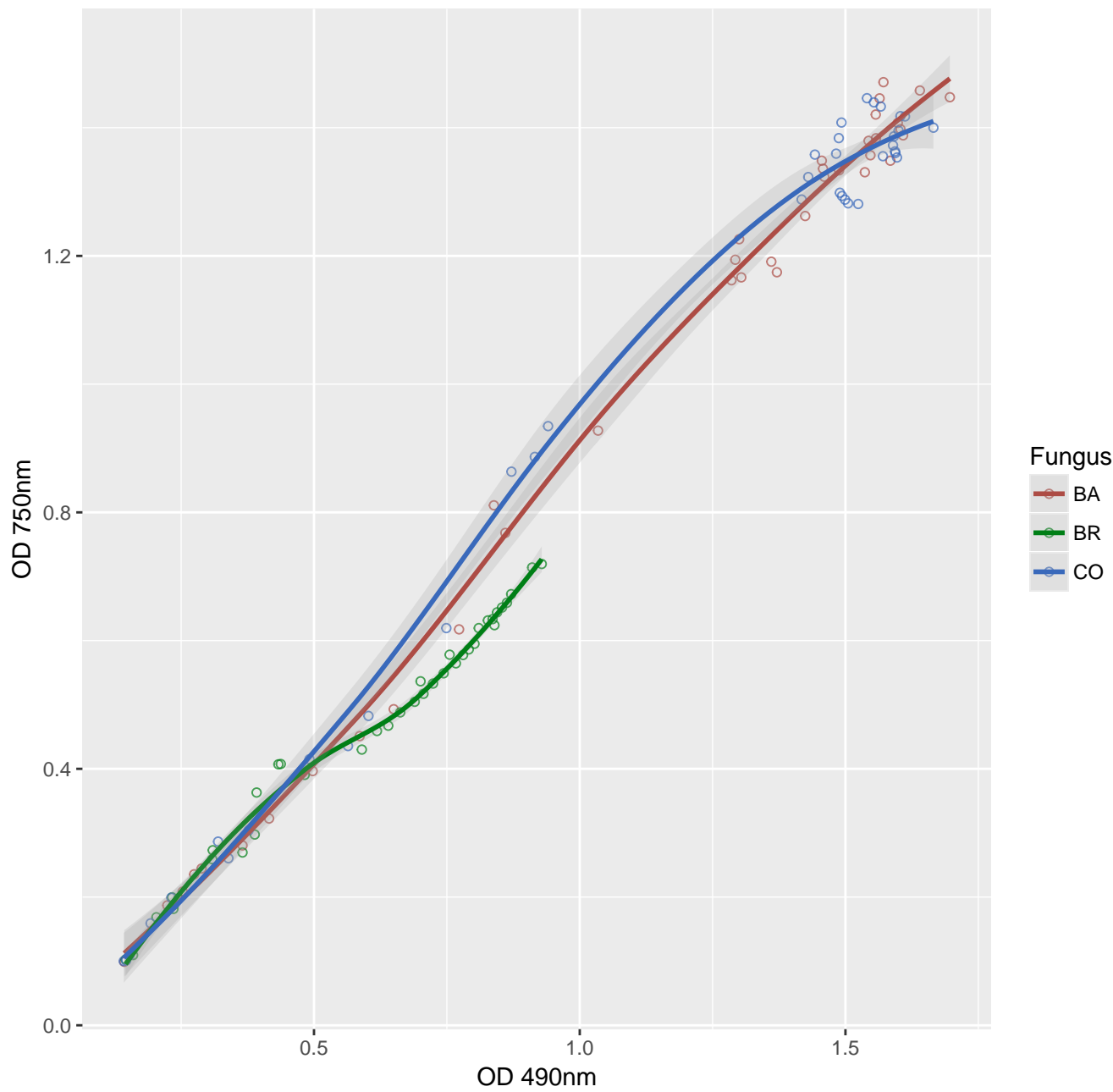
D.Melibiose



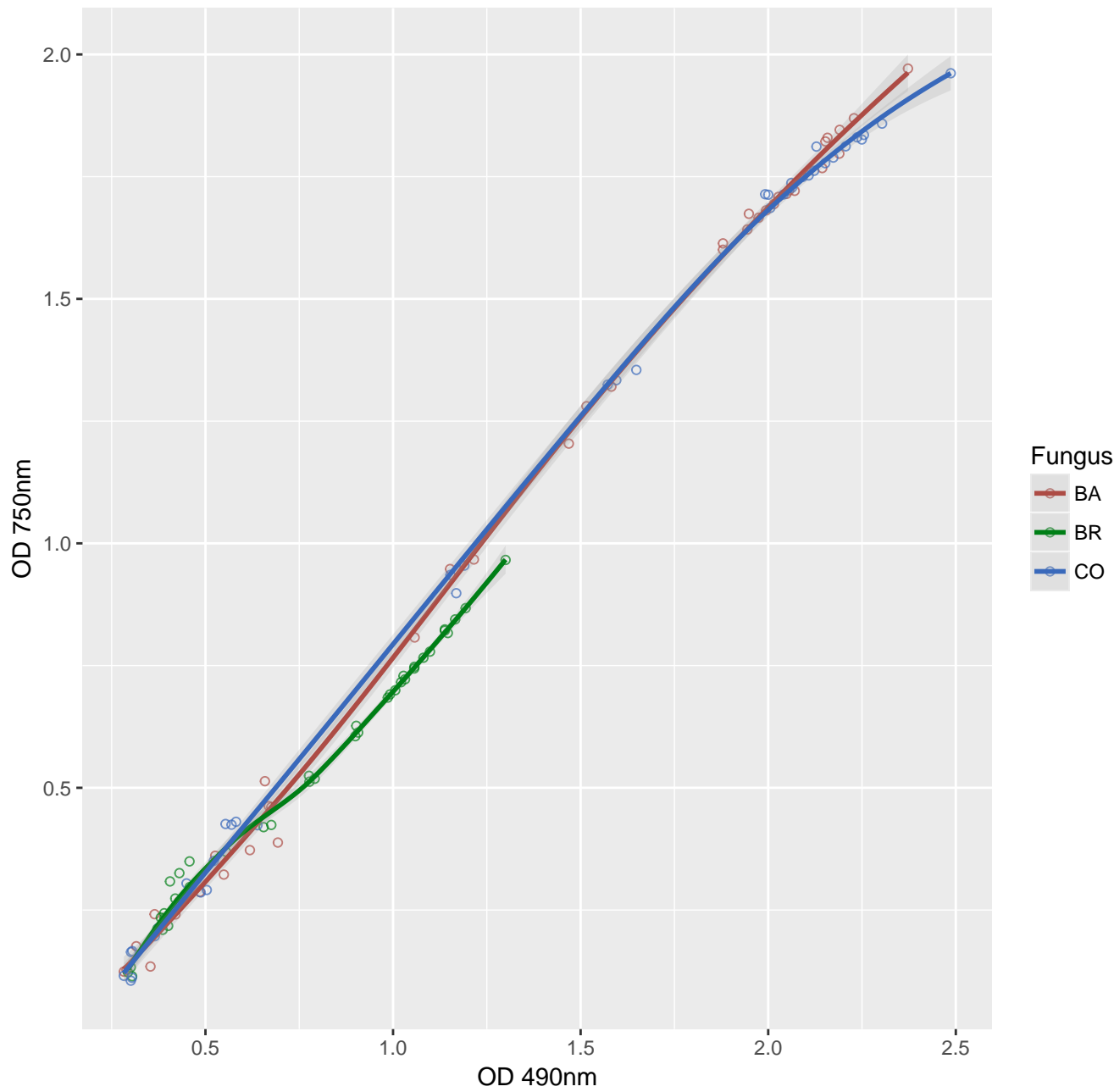
D.Psicose



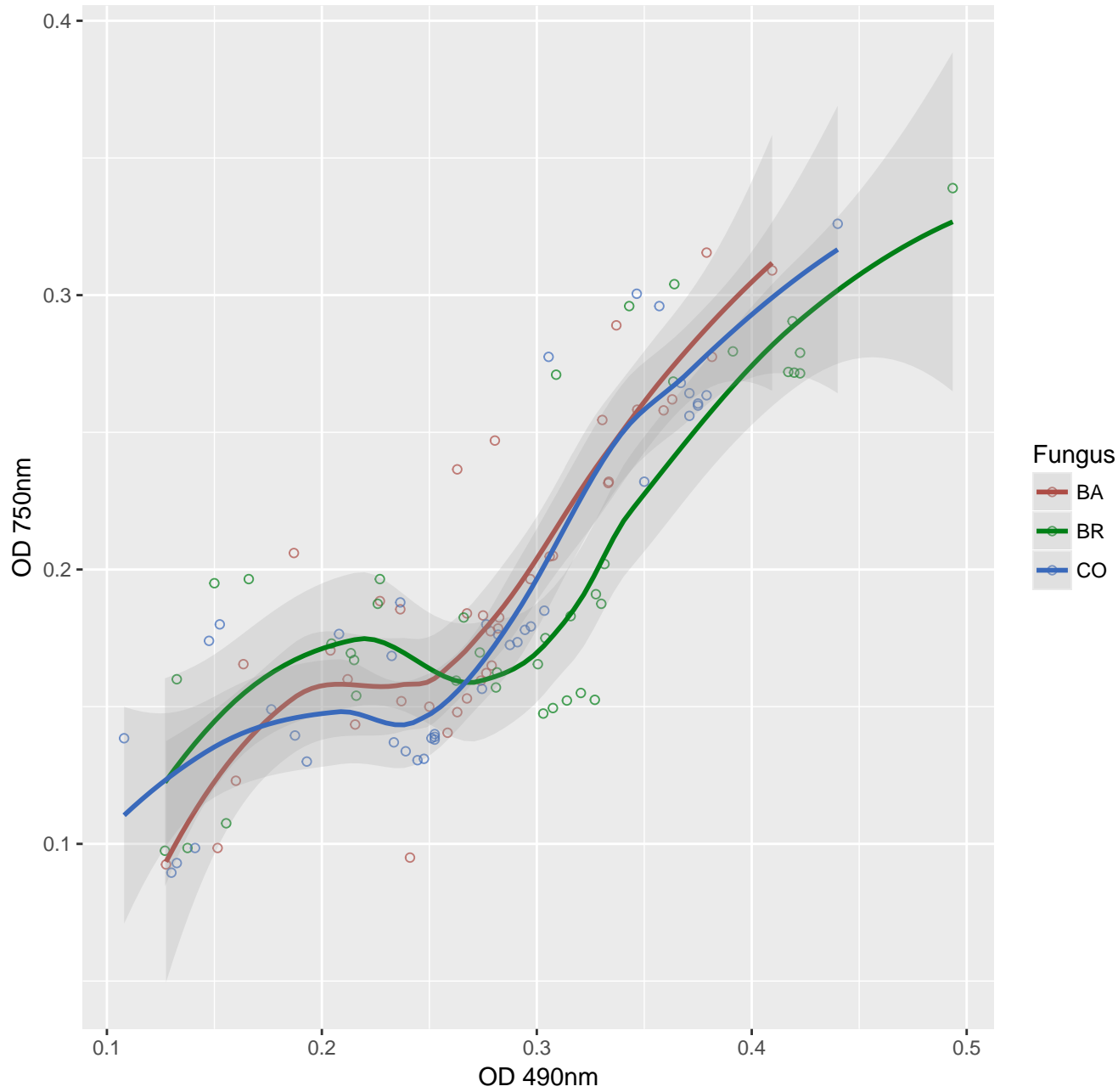
D.Raffinose



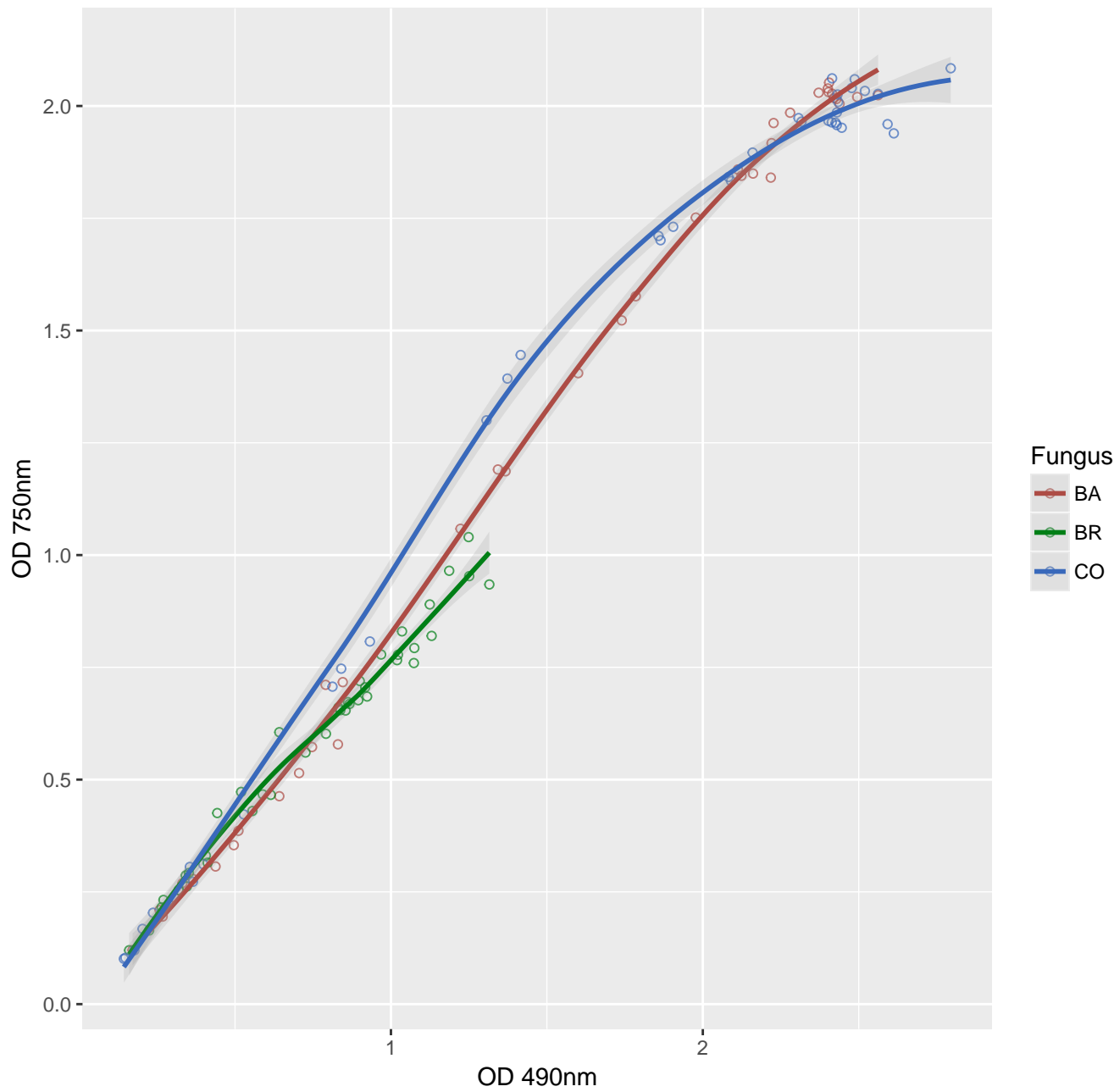
D.Ribose



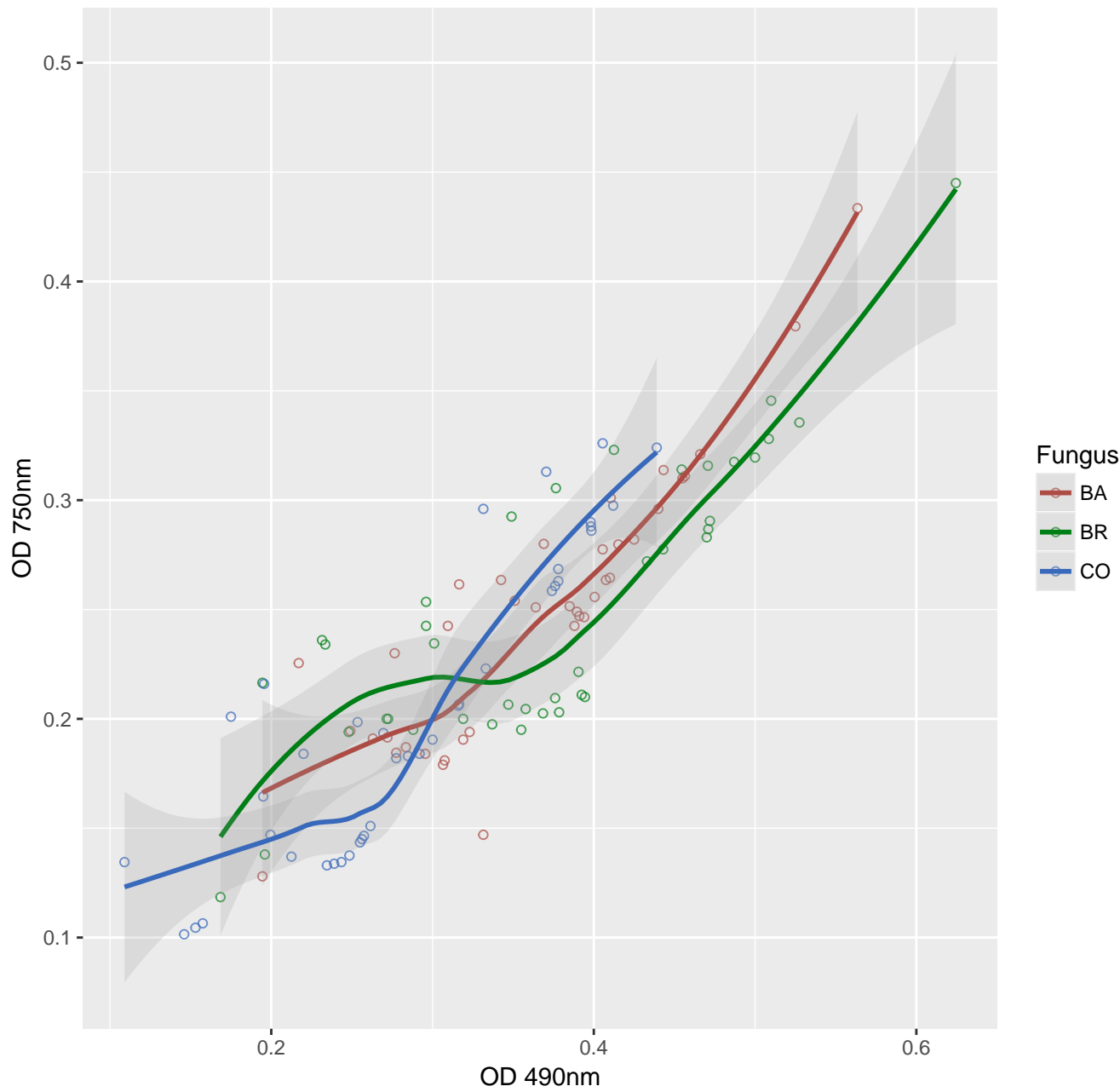
D.SaccharinAcid



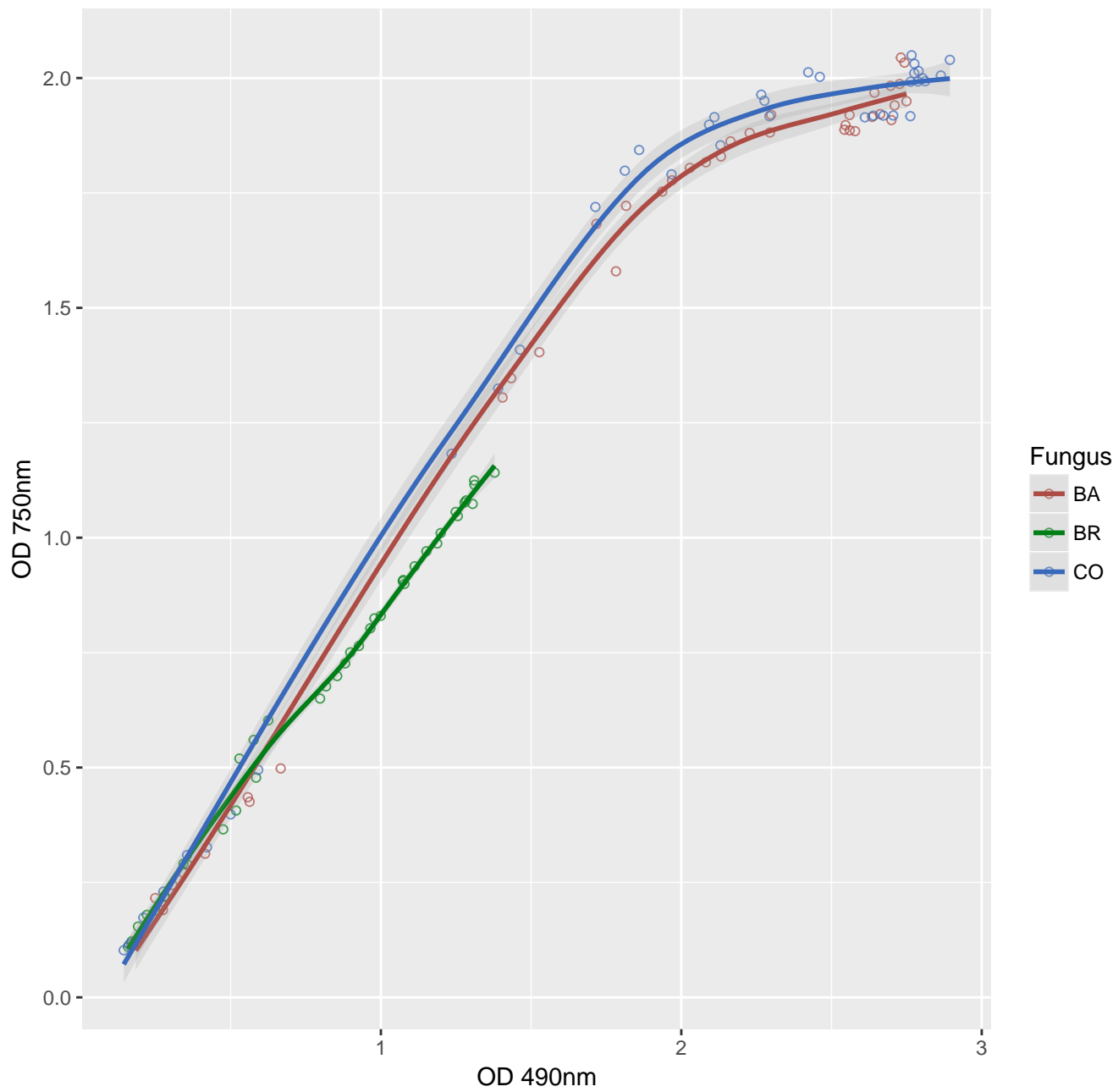
D.Sorbitol



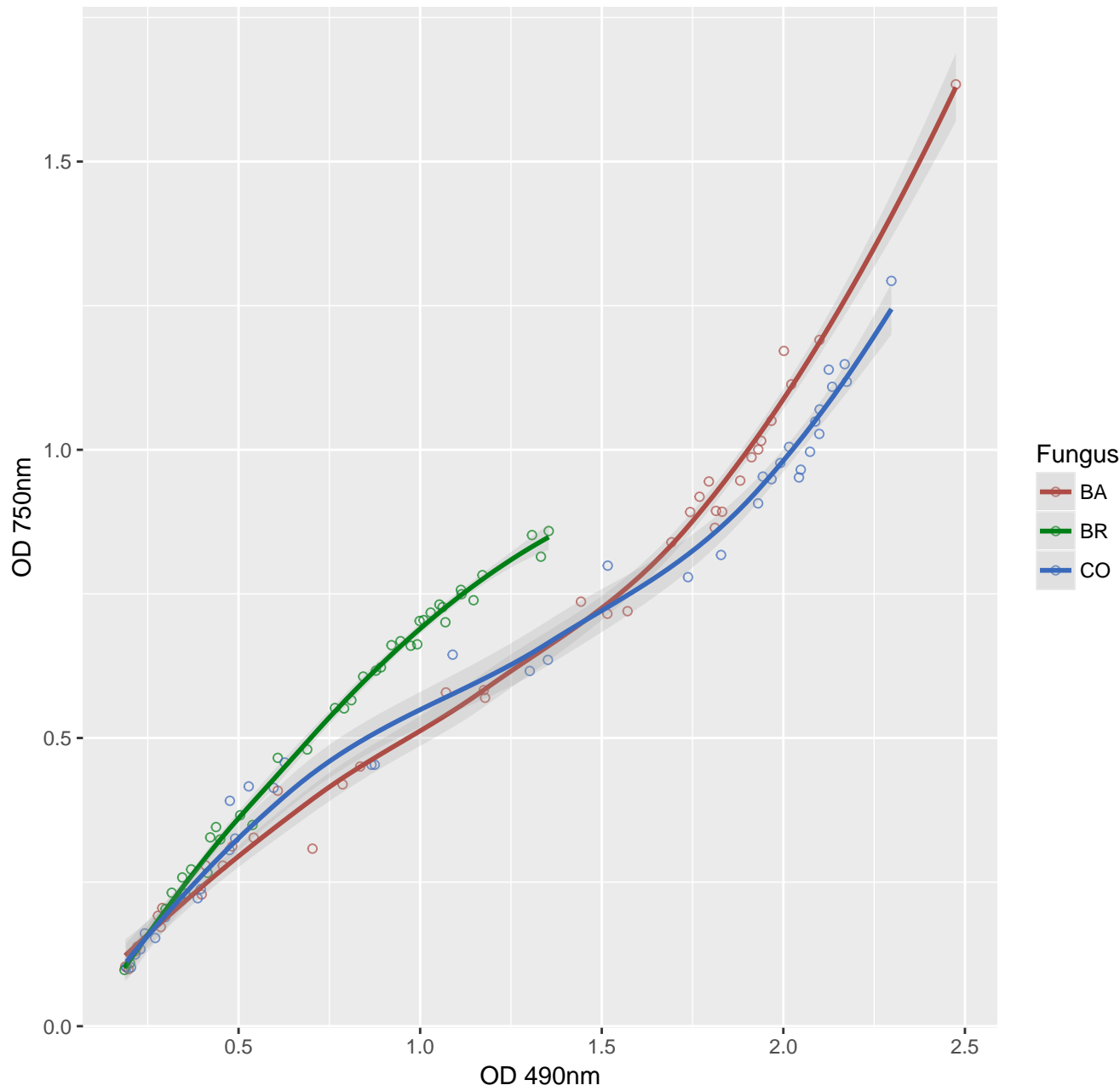
D.Tagatose



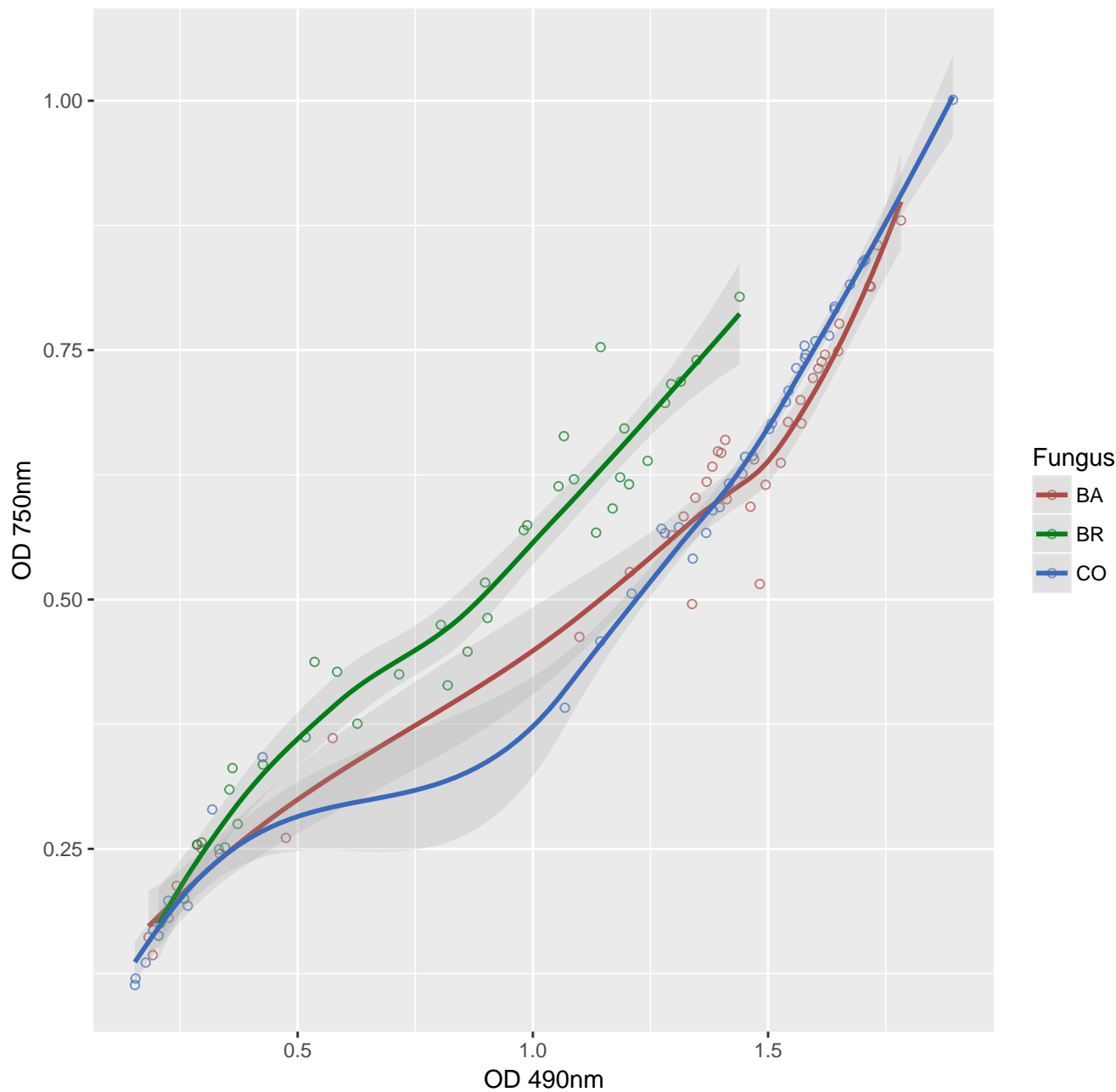
D.Trehalose



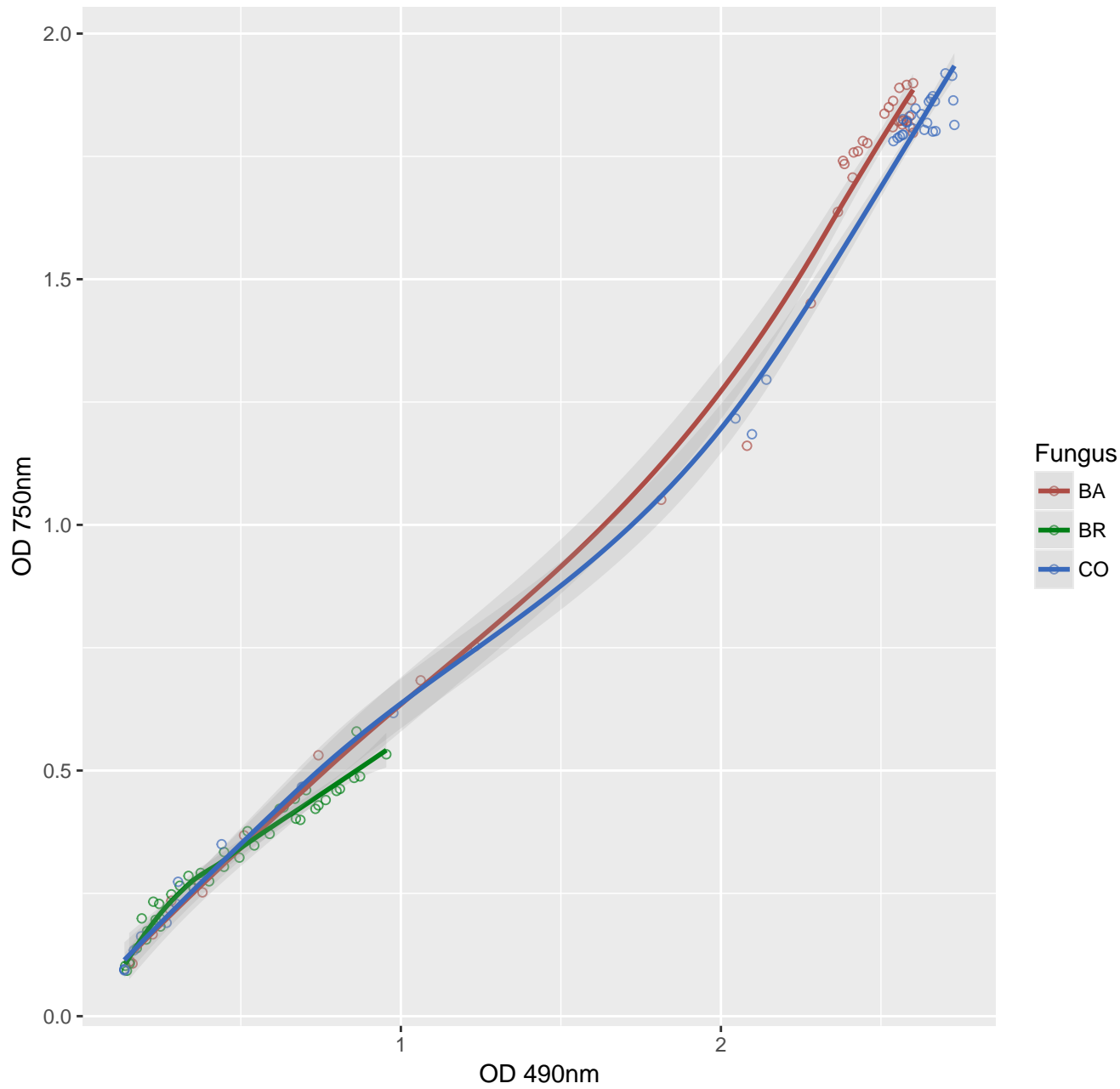
D.Xylose



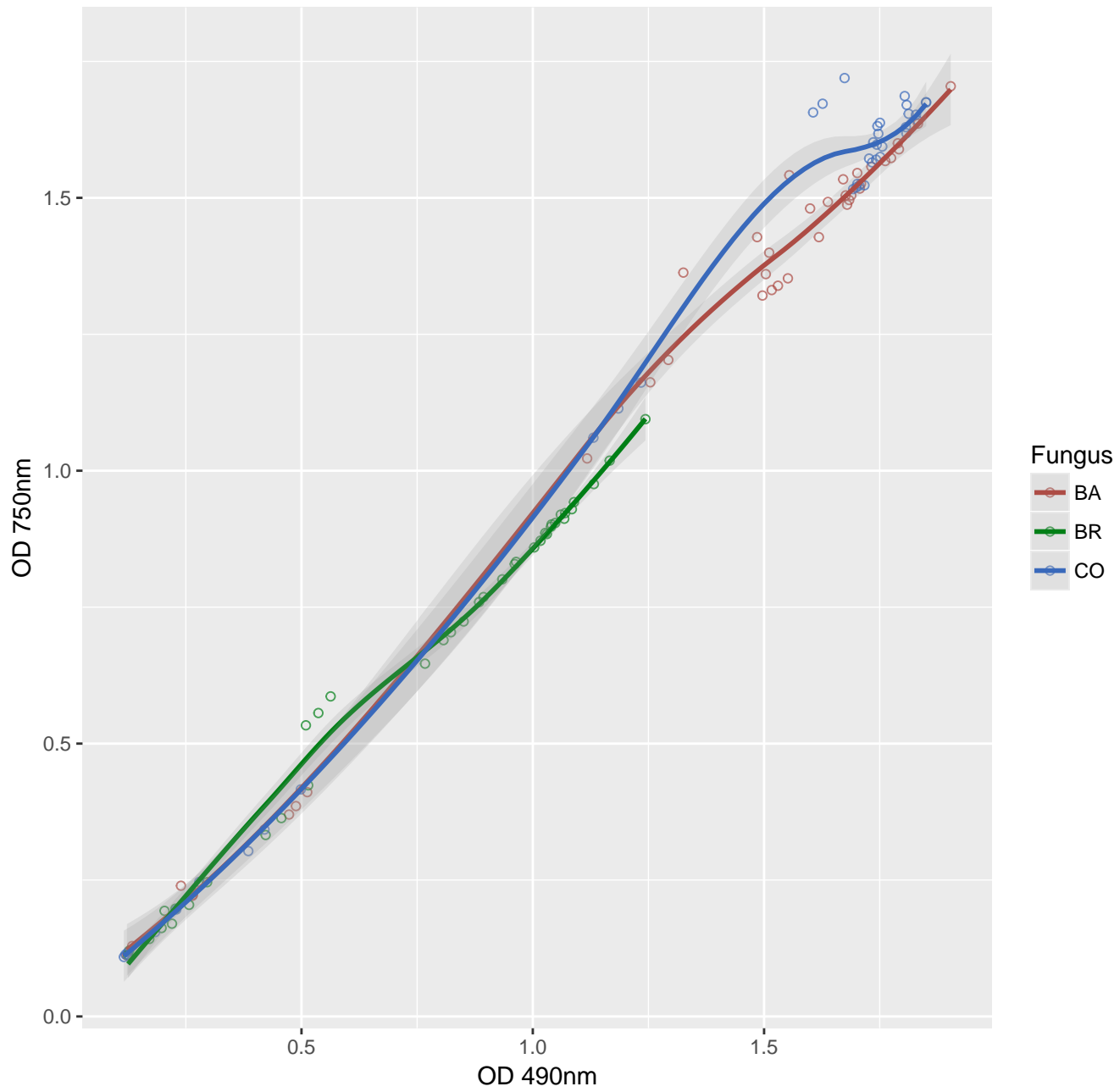
FumaricAcid



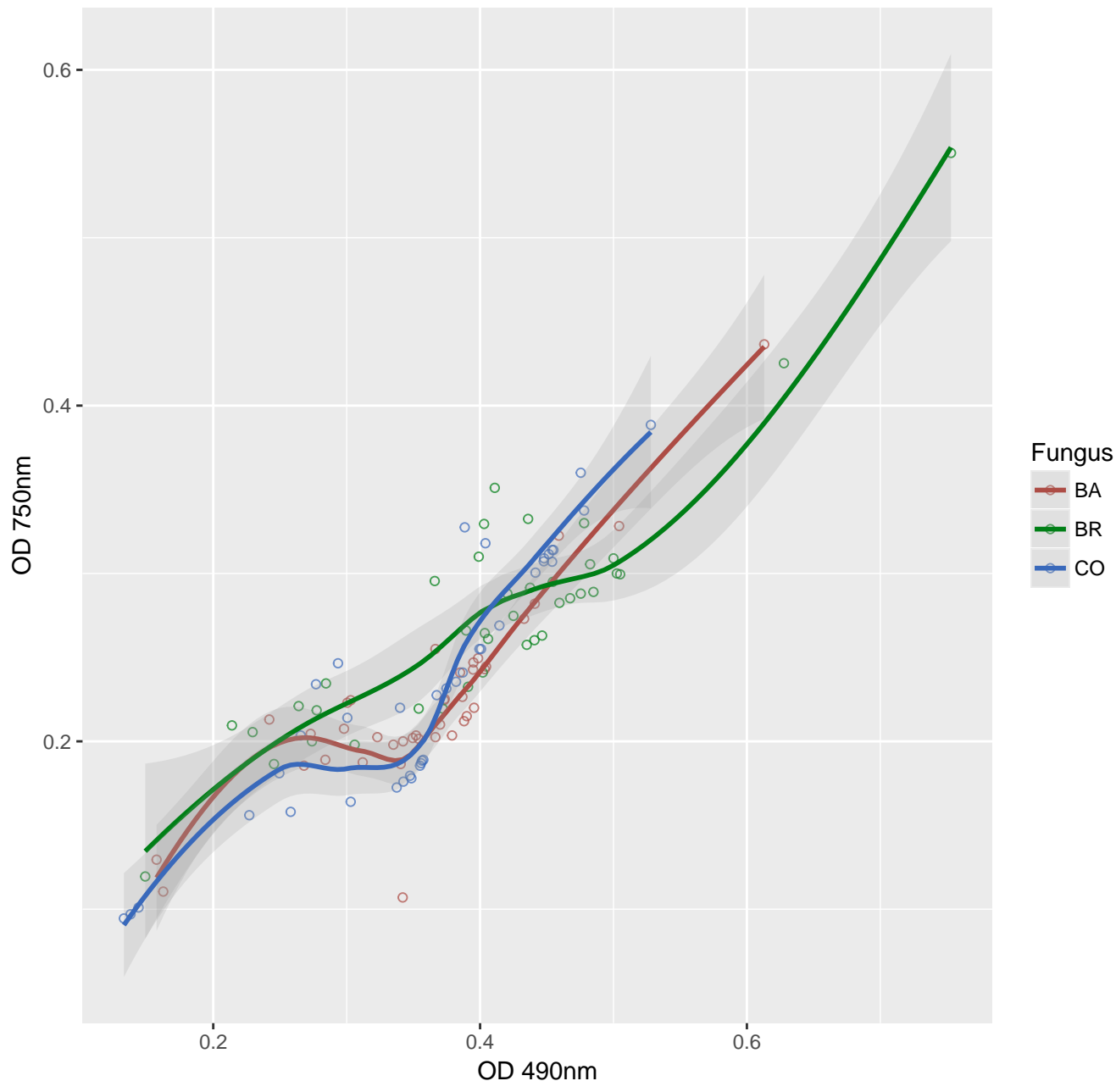
G.Amino.butyricAcid



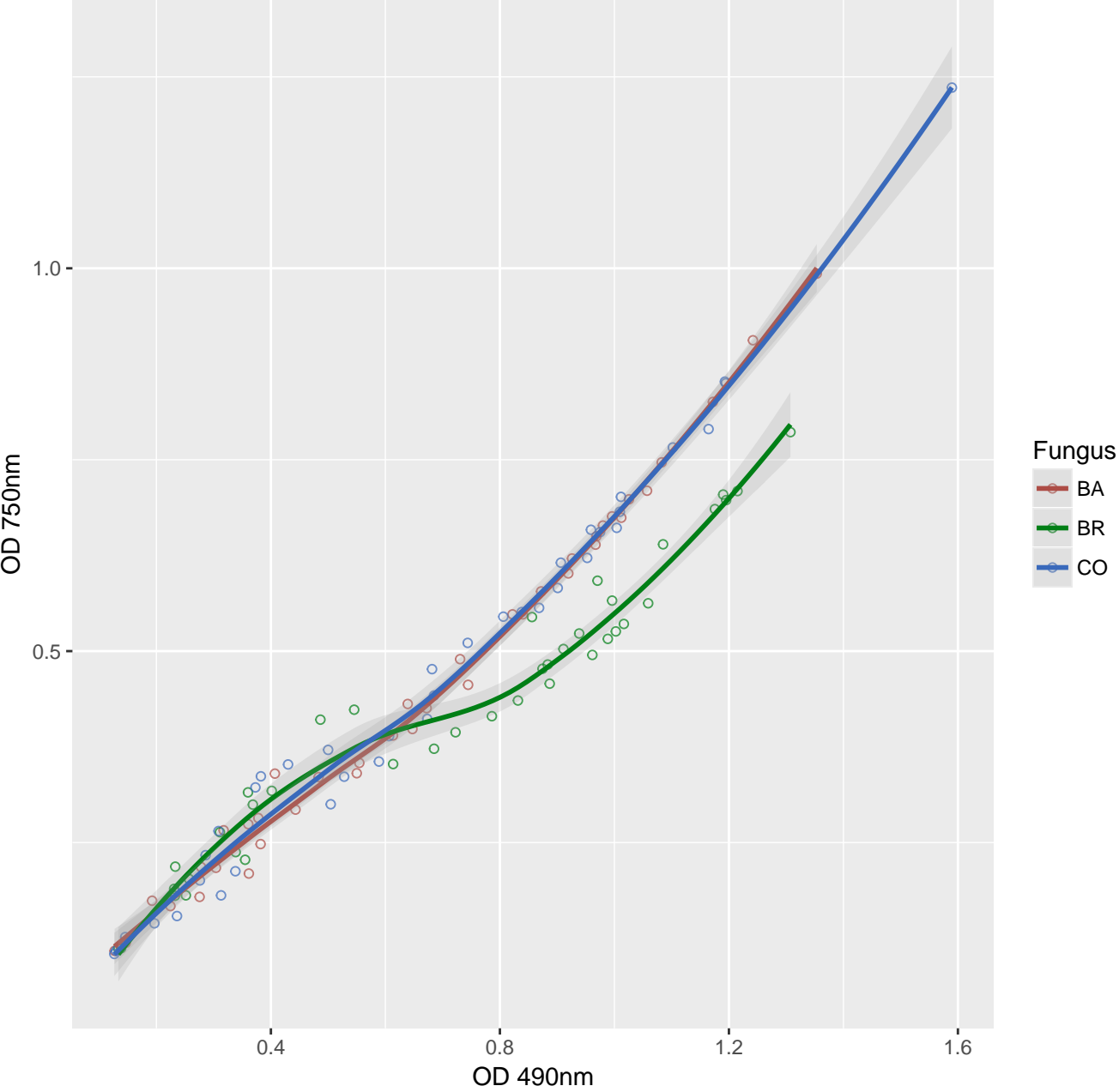
Gentibiose



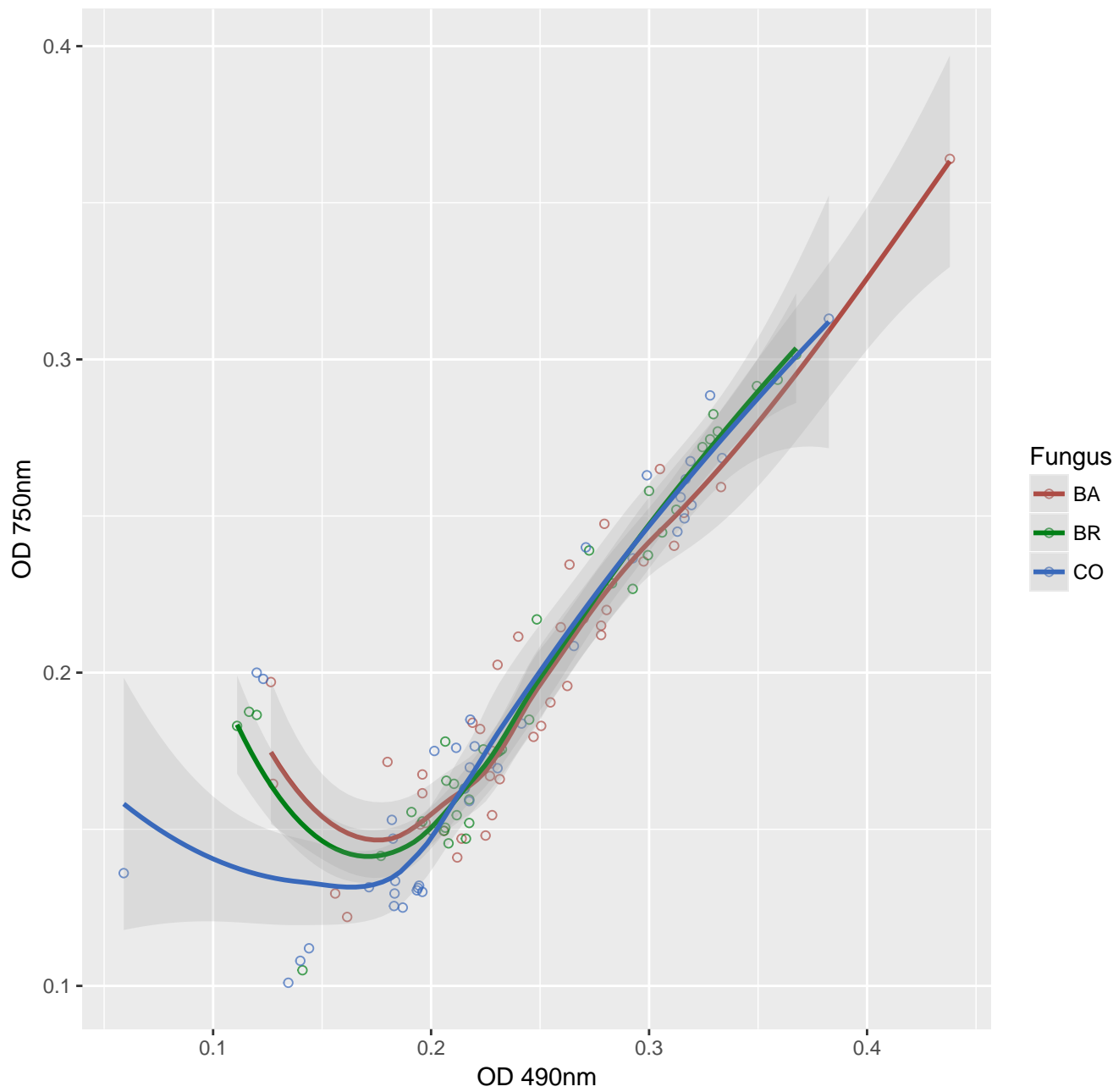
G.Hydroxy.butyricAcid



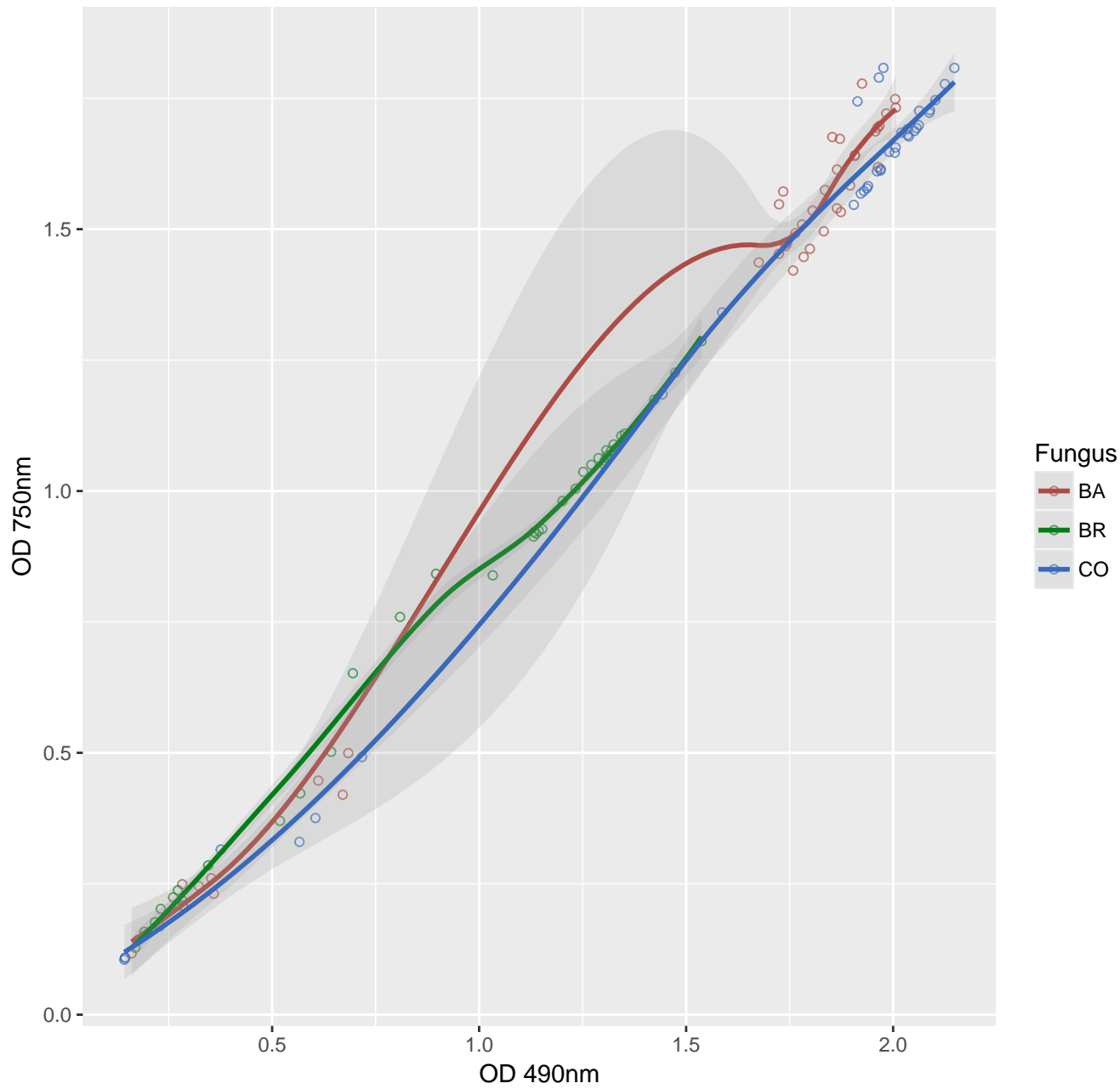
Glucose.1.Phosphate



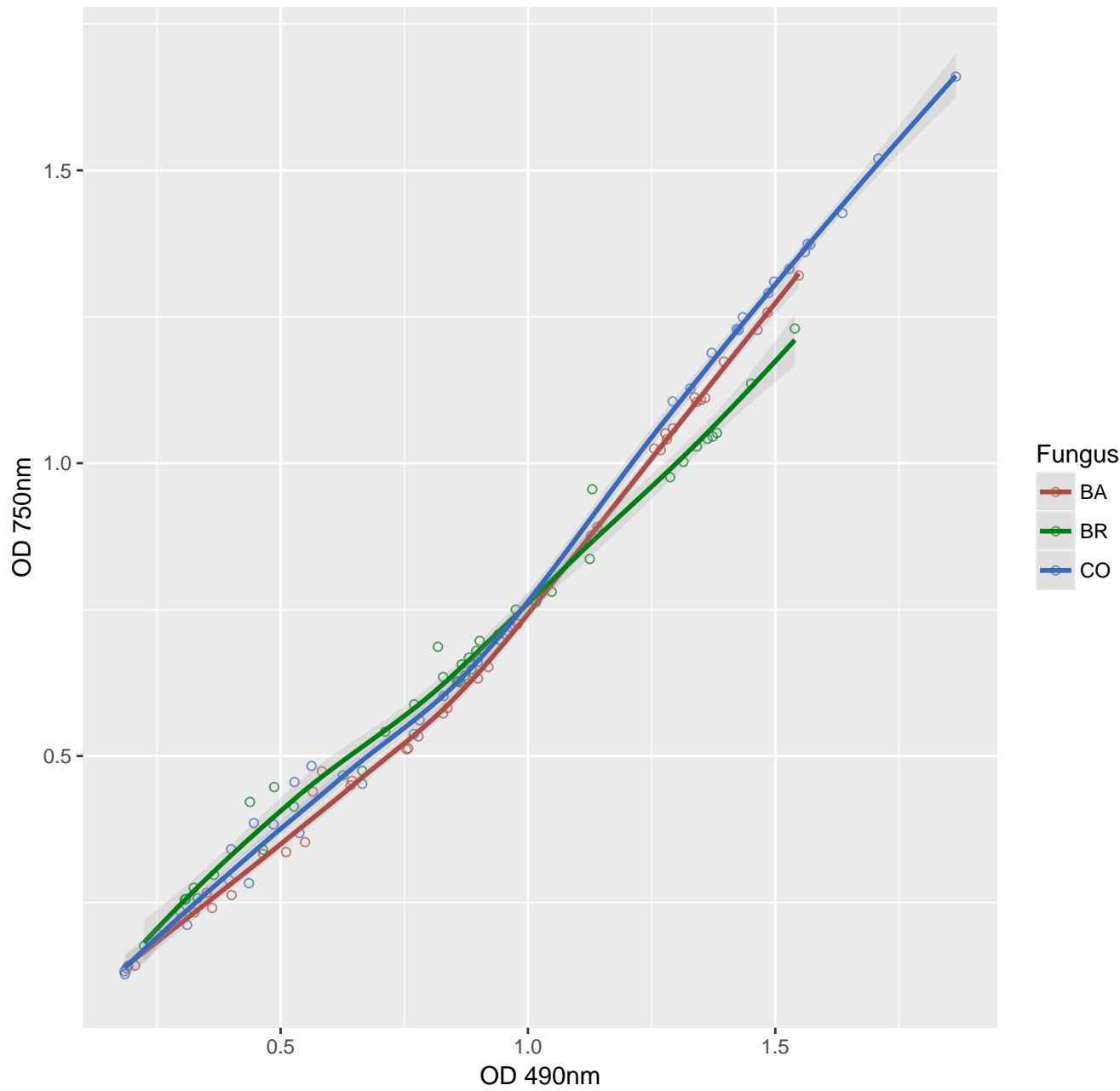
Glucuronamide



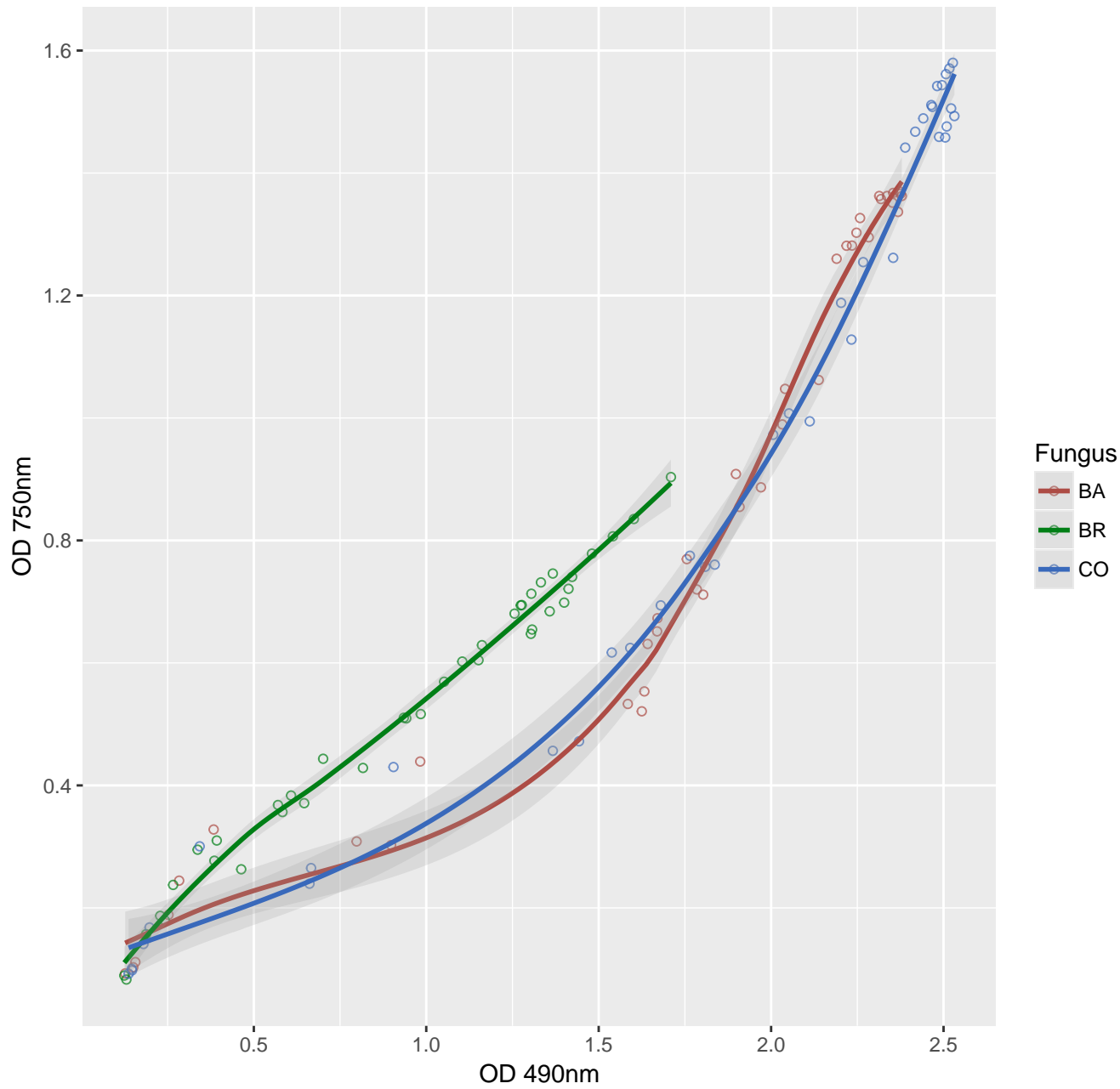
Glycerol



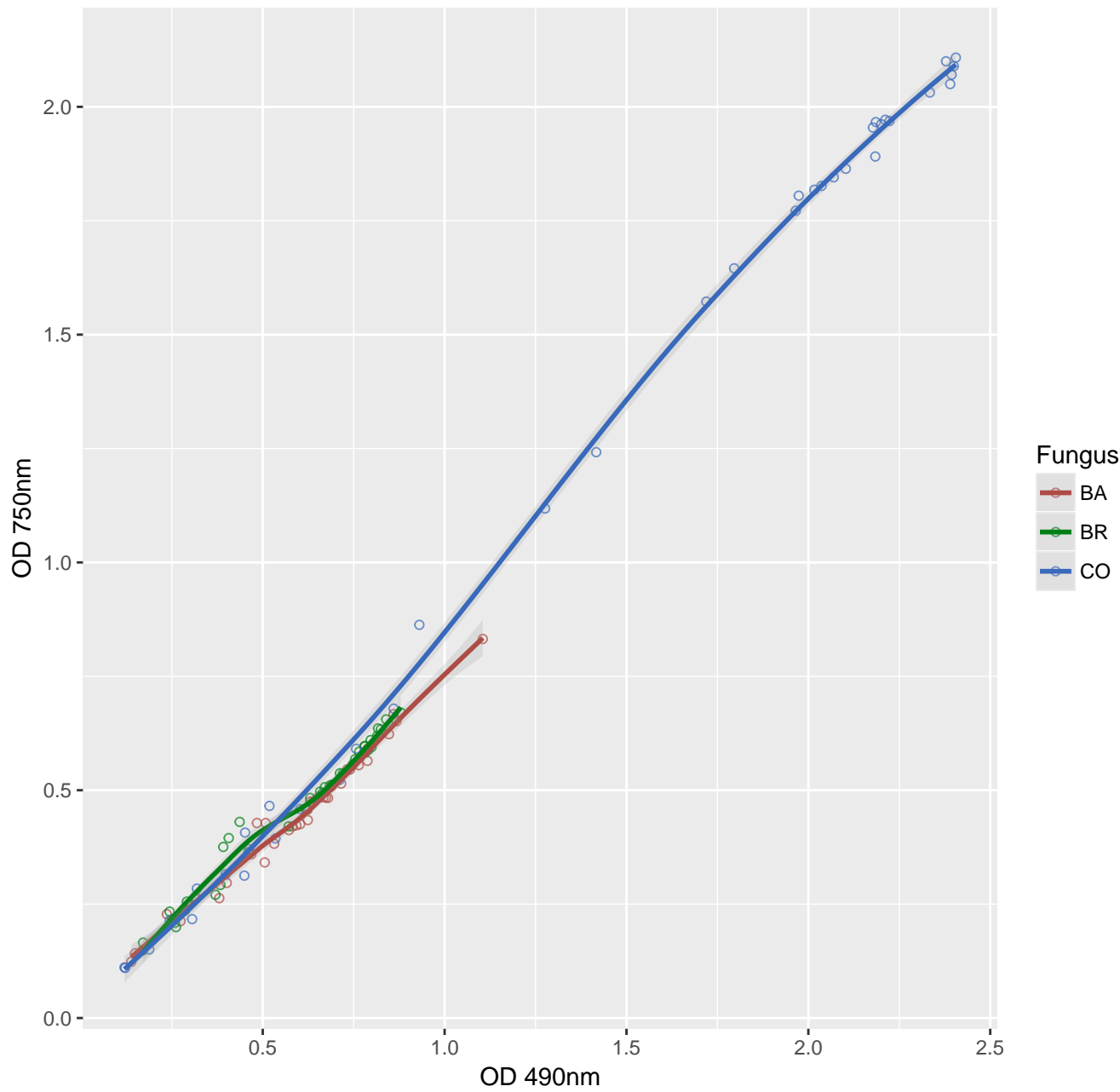
Glycogen



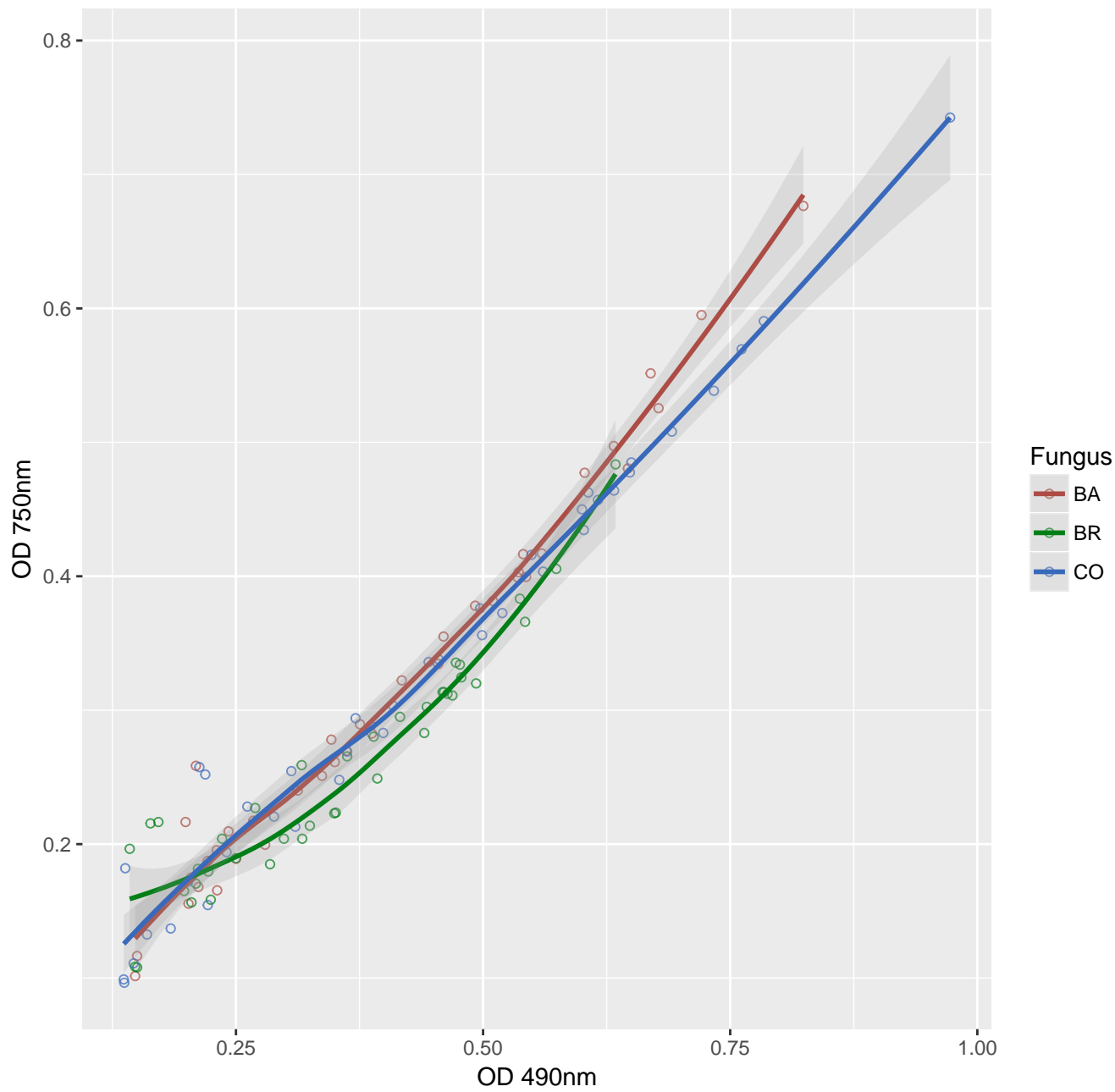
Glycyl.L.GlutamicAcid



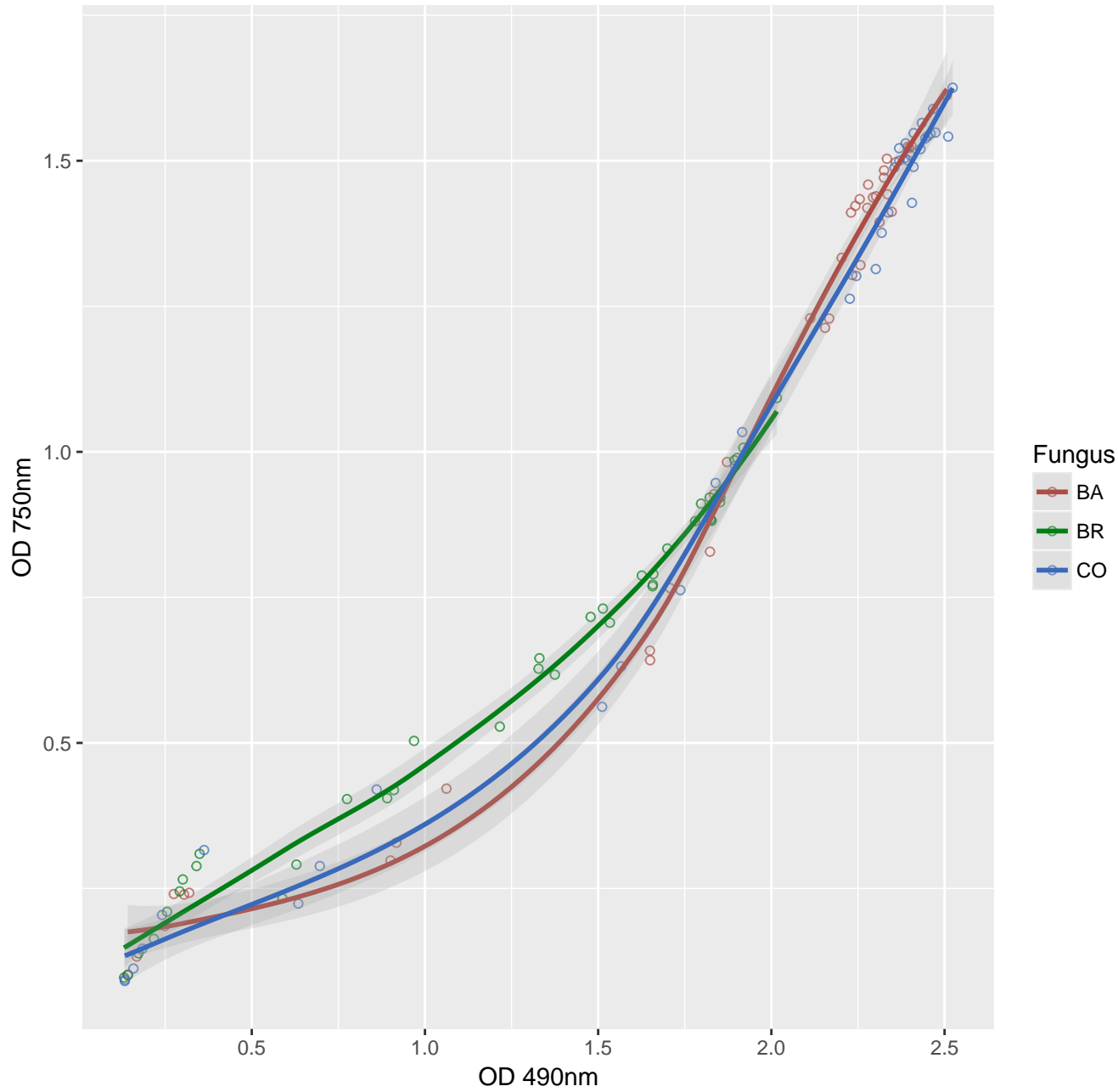
I.Erythritol



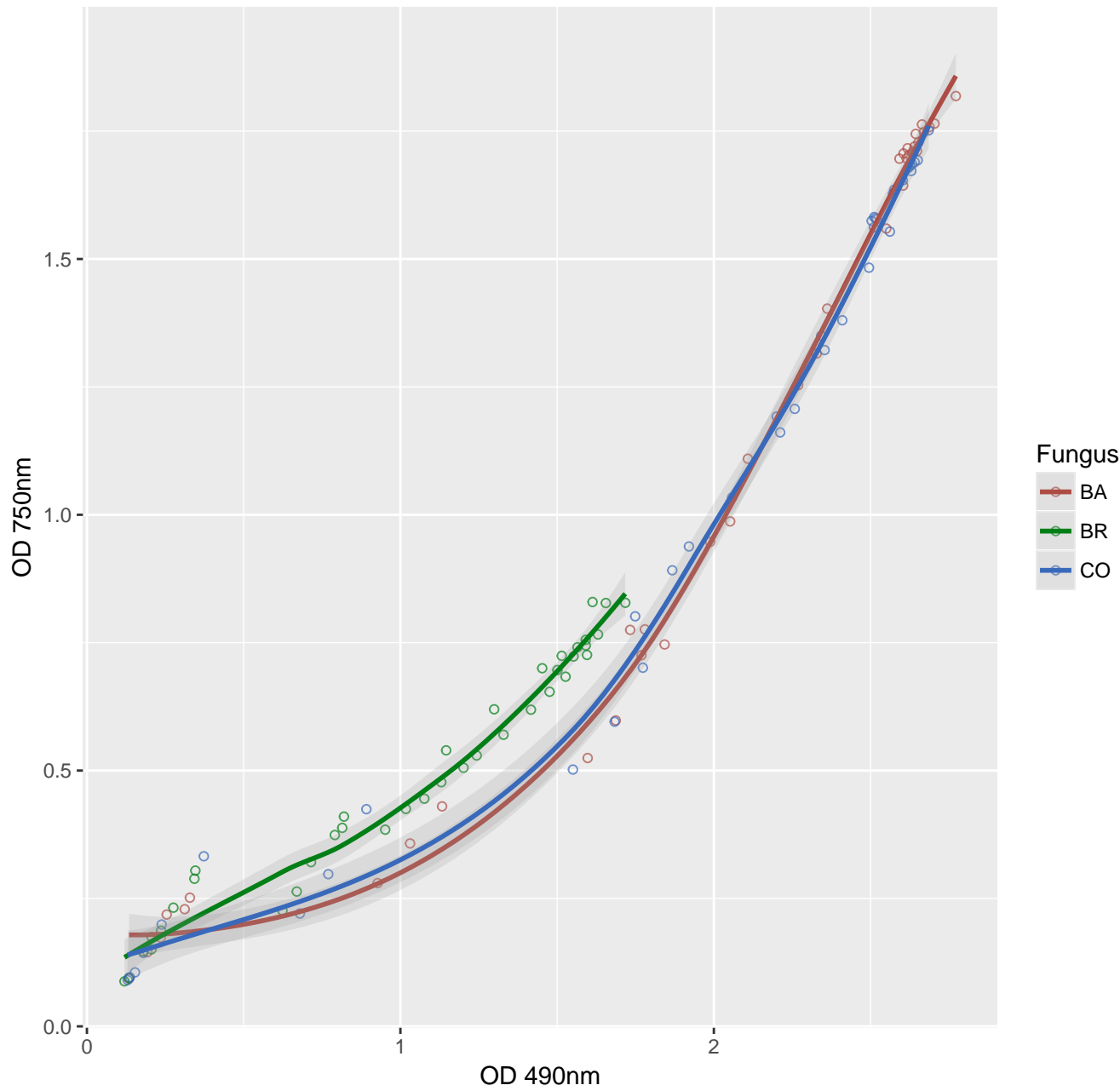
Lactulose



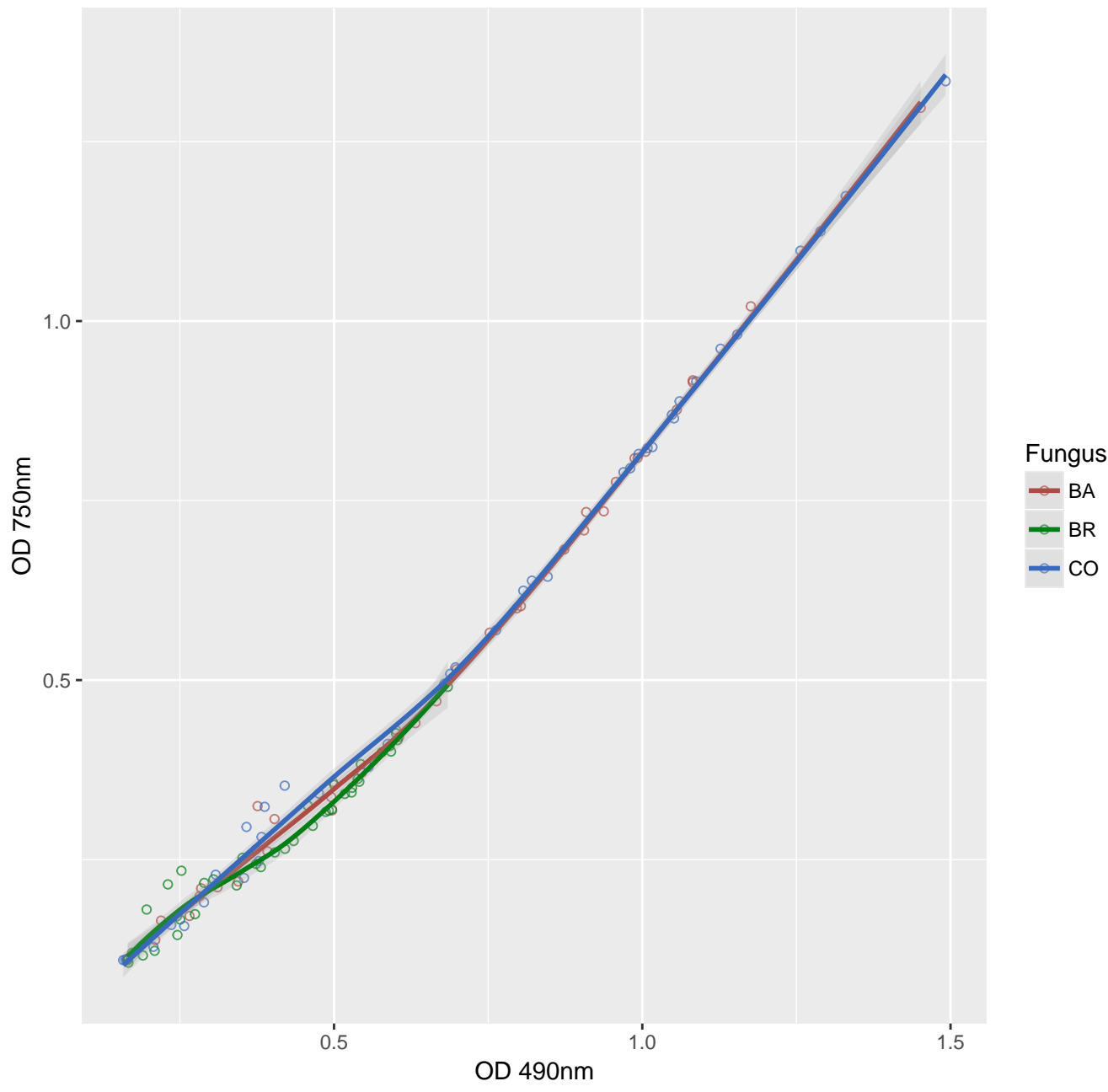
L.Alanine



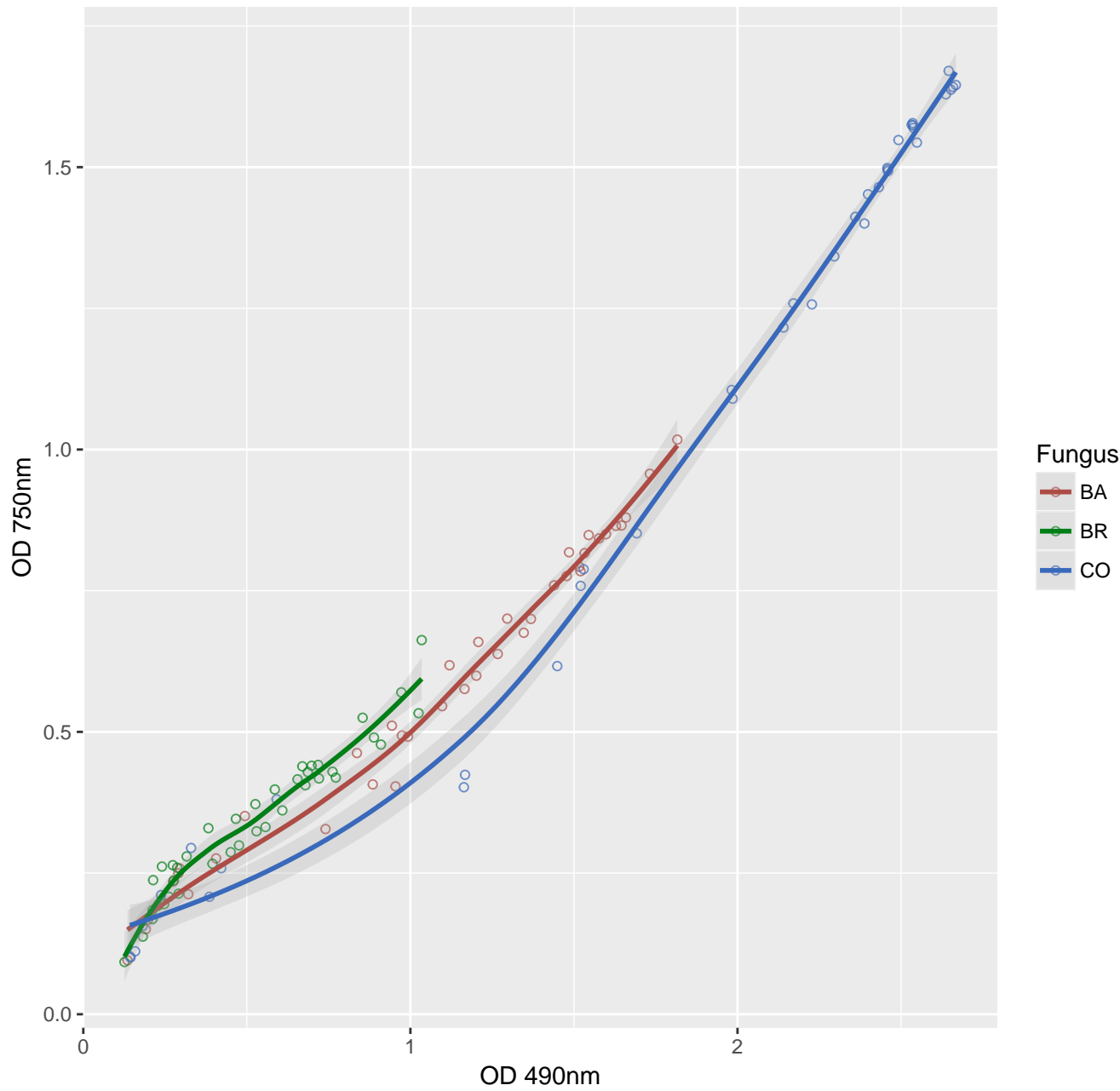
L.Alanyl.Glycine



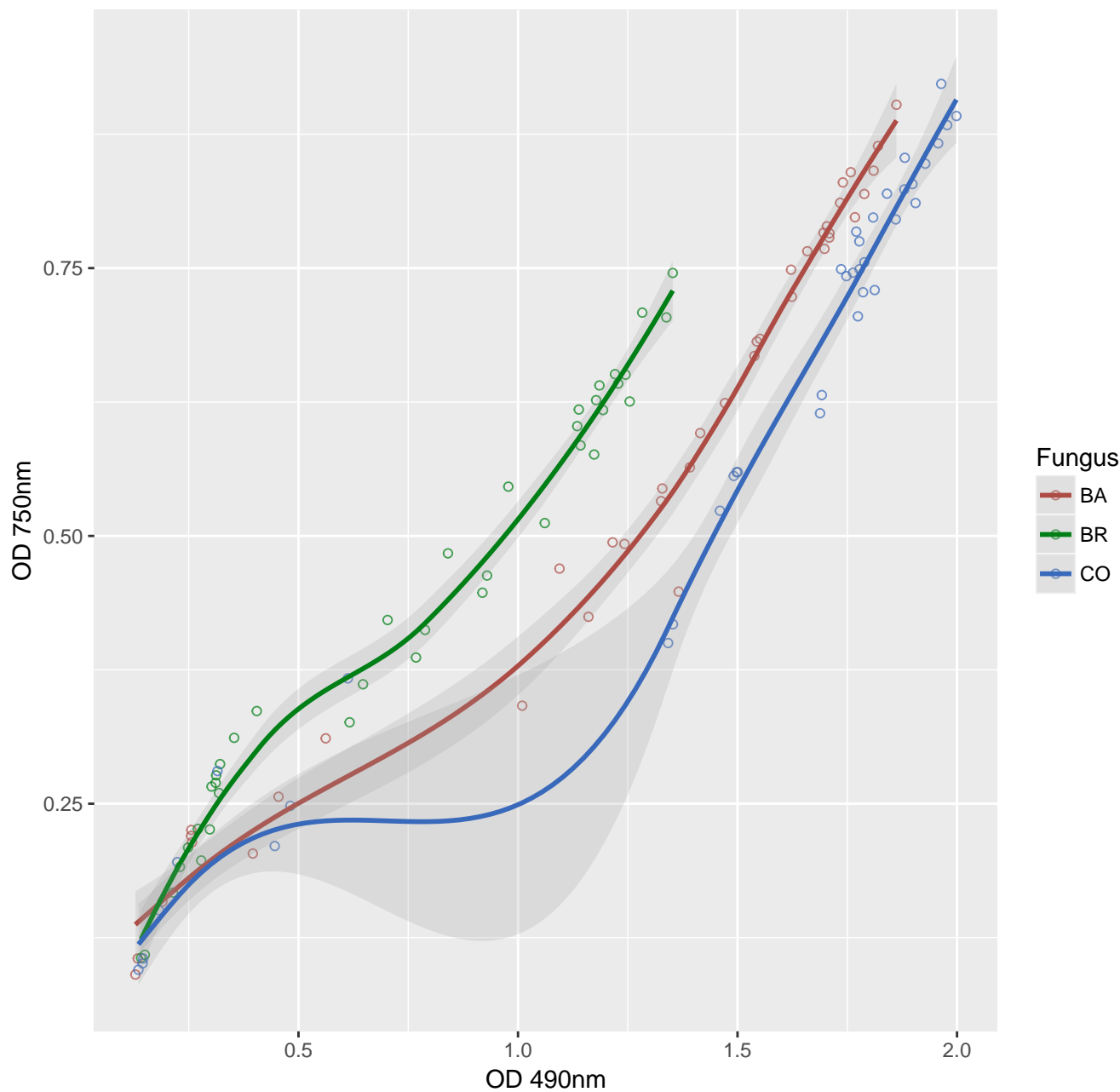
L.Arabinose



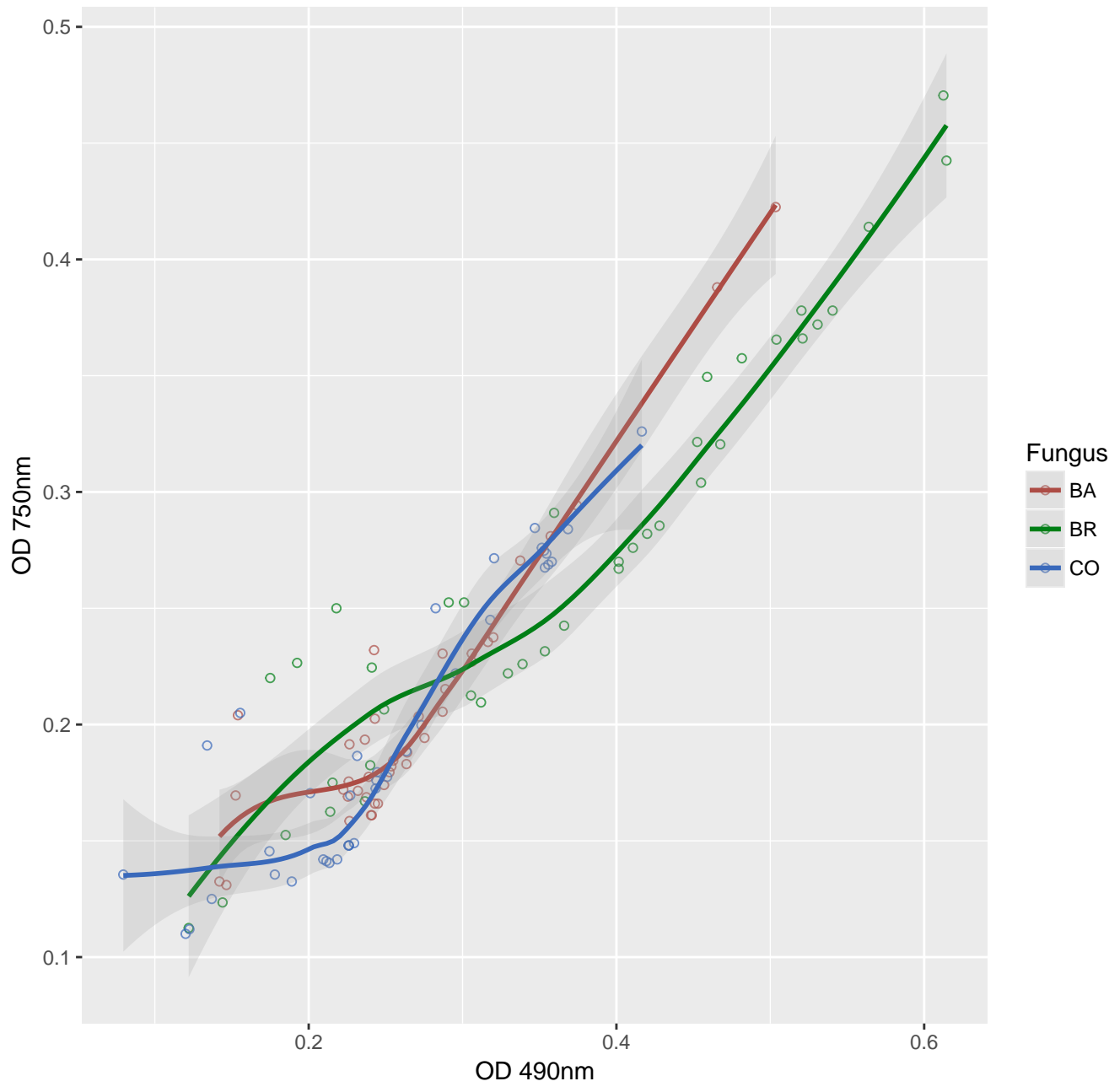
L.Asparagine



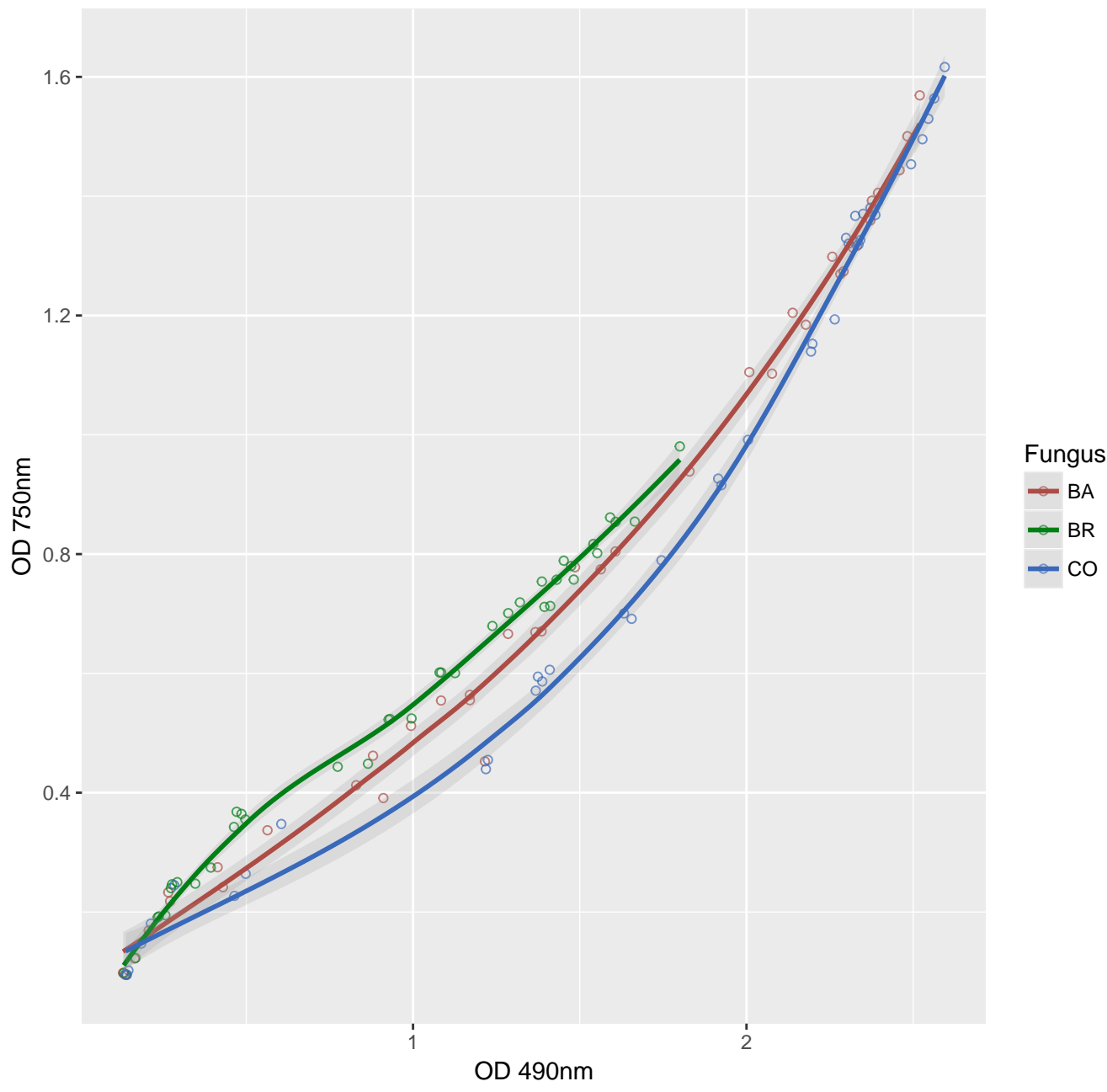
L.AsparticAcid



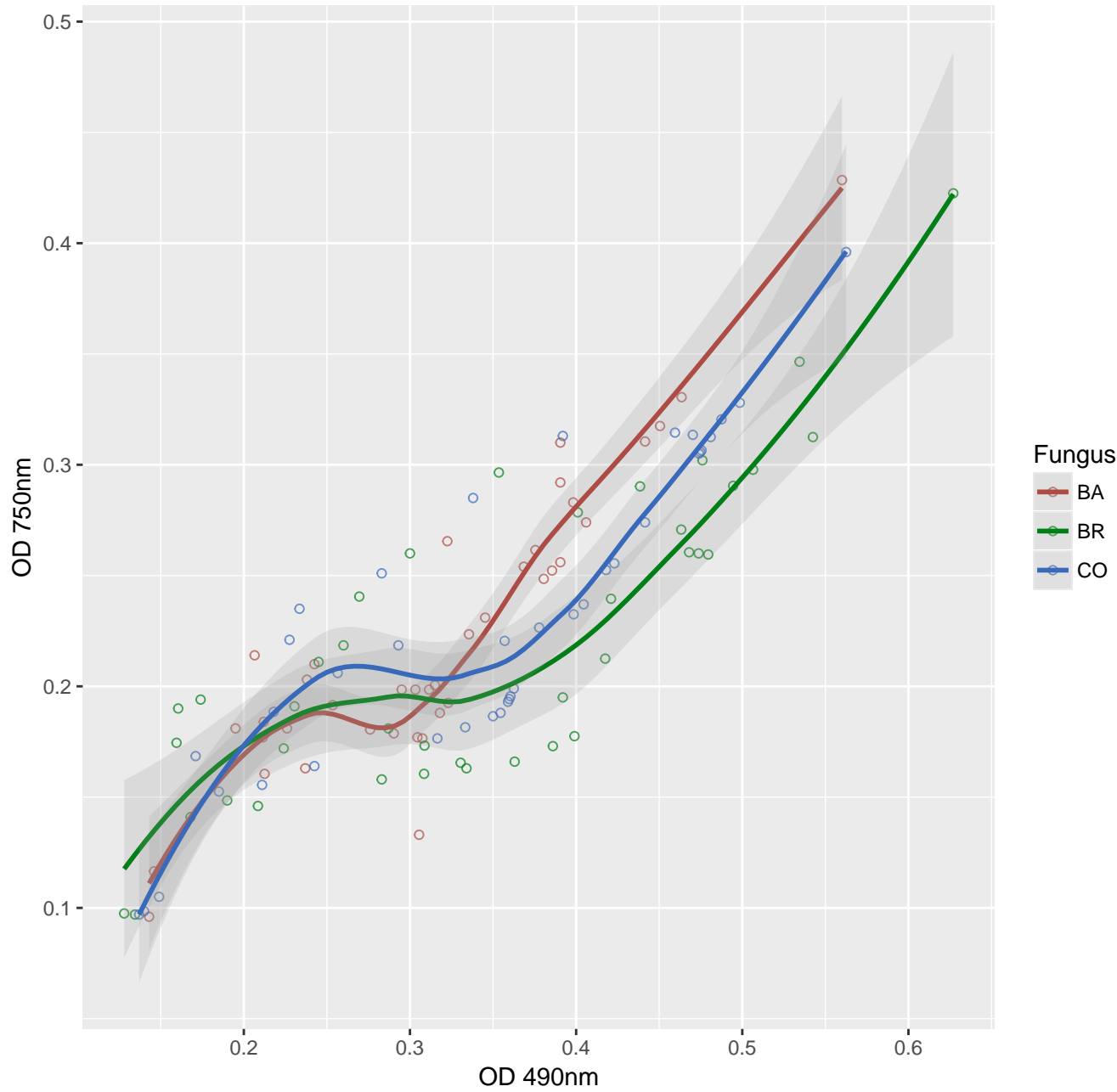
L.Fucose



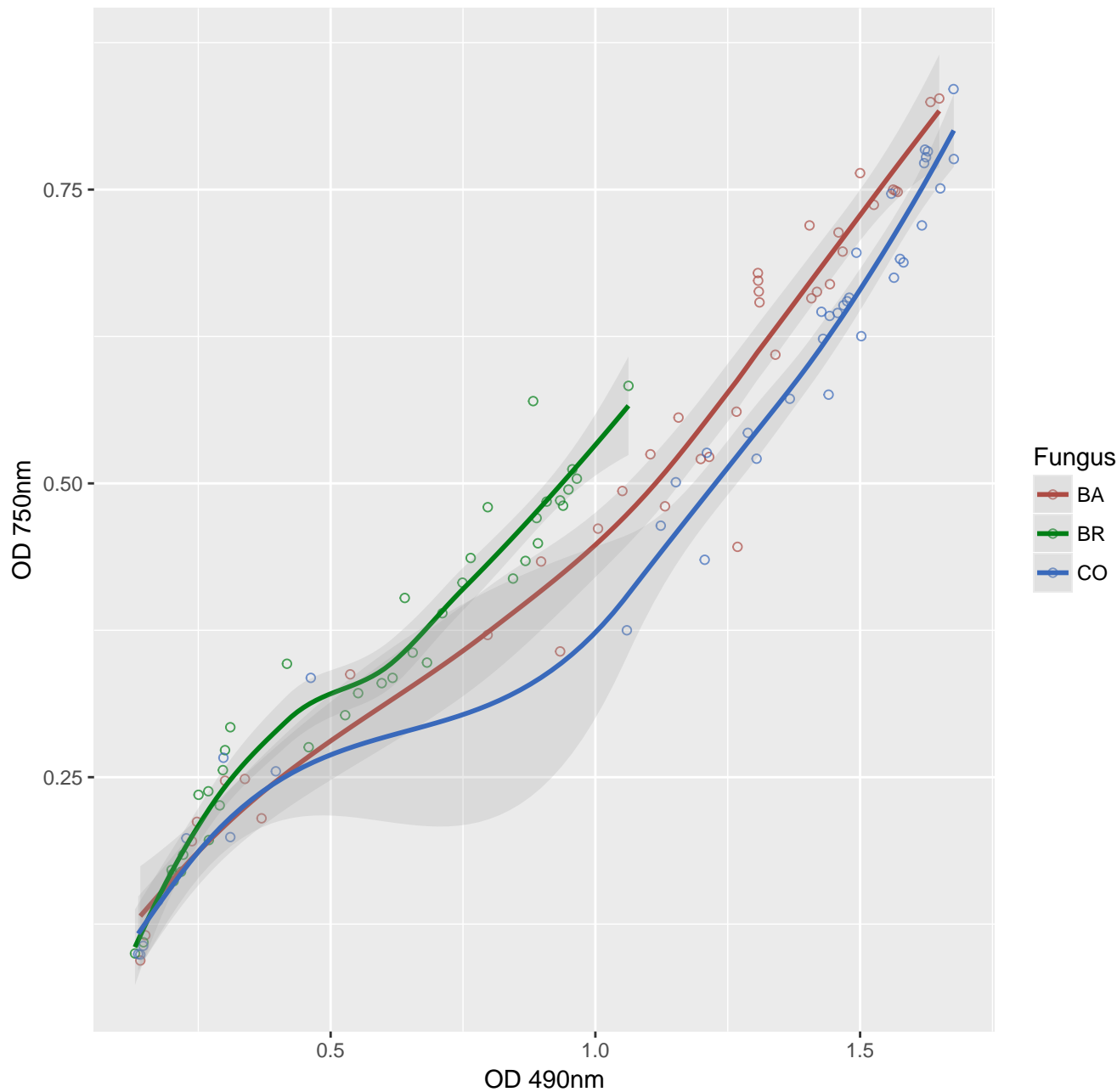
L.GlutamicAcid



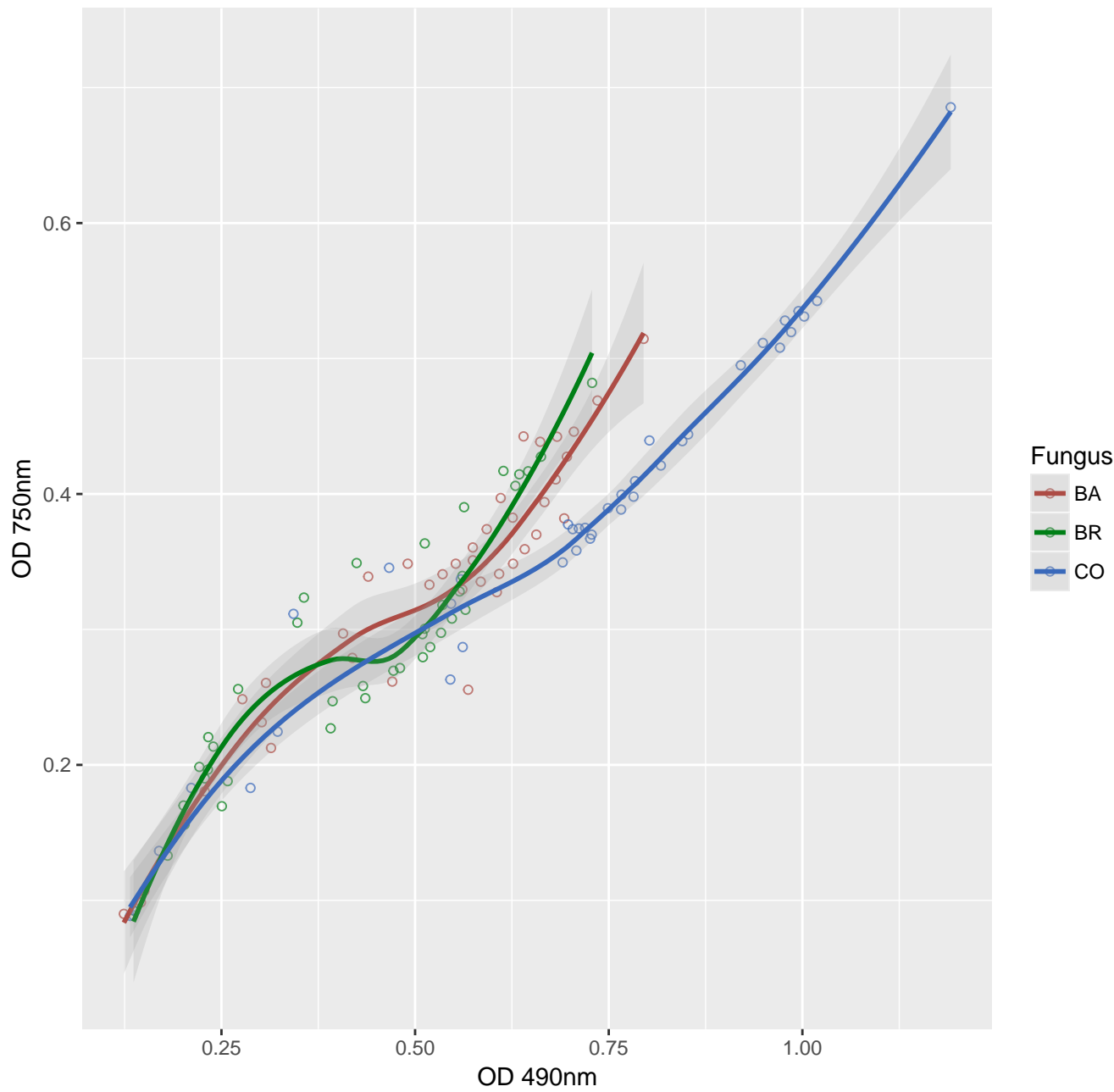
L.LacticAcid



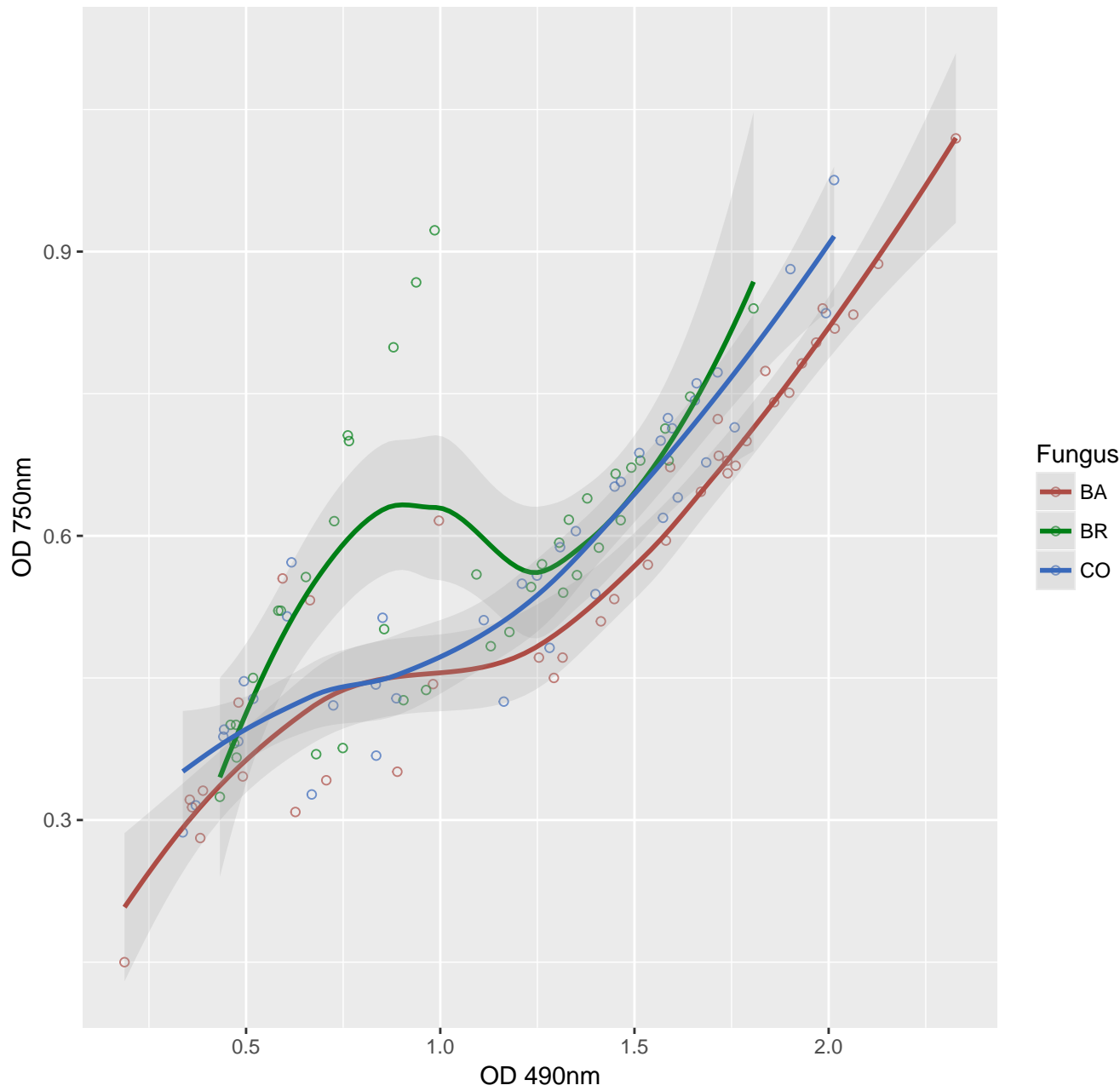
L.MalicAcid



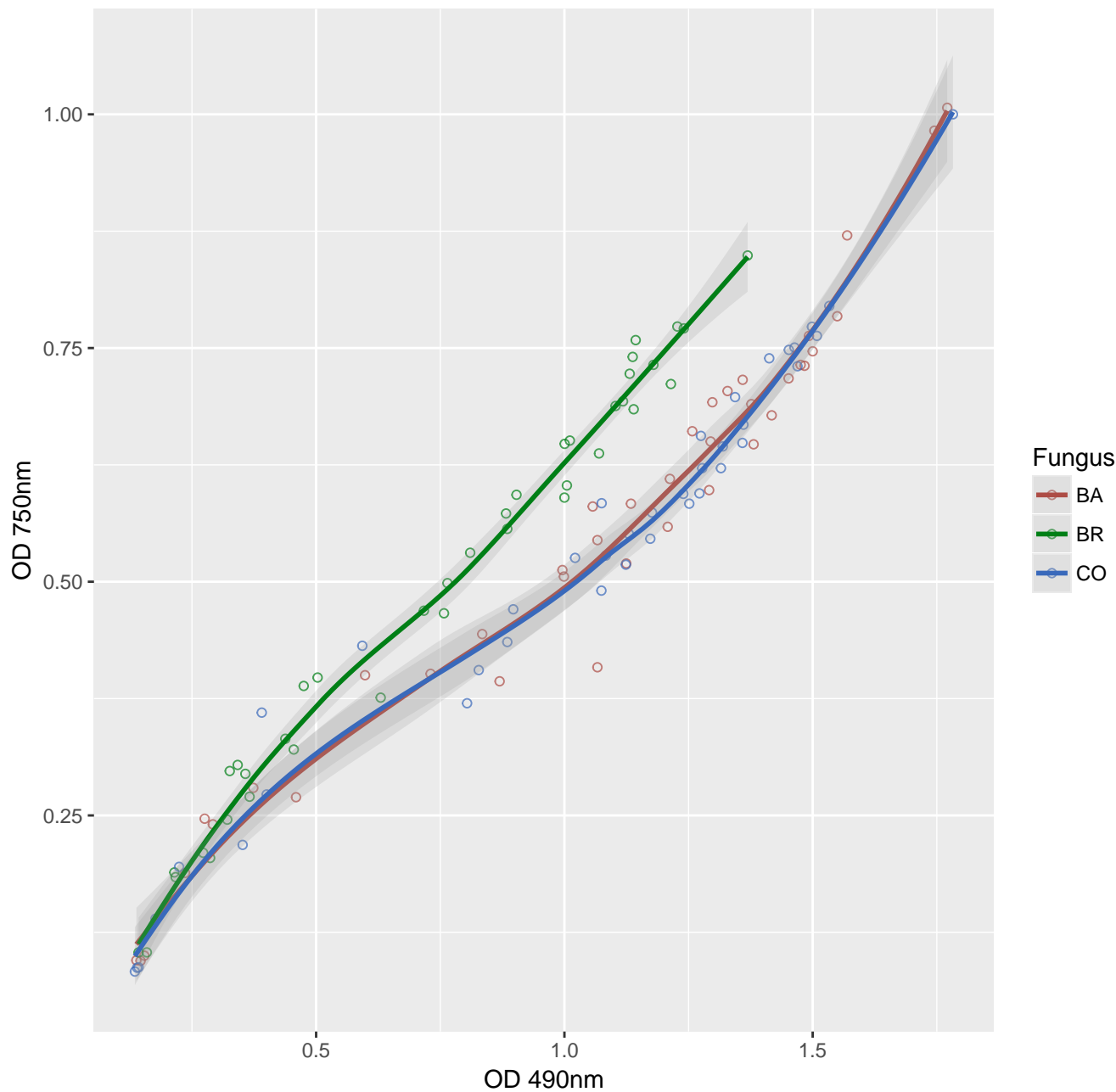
L.Ornithine



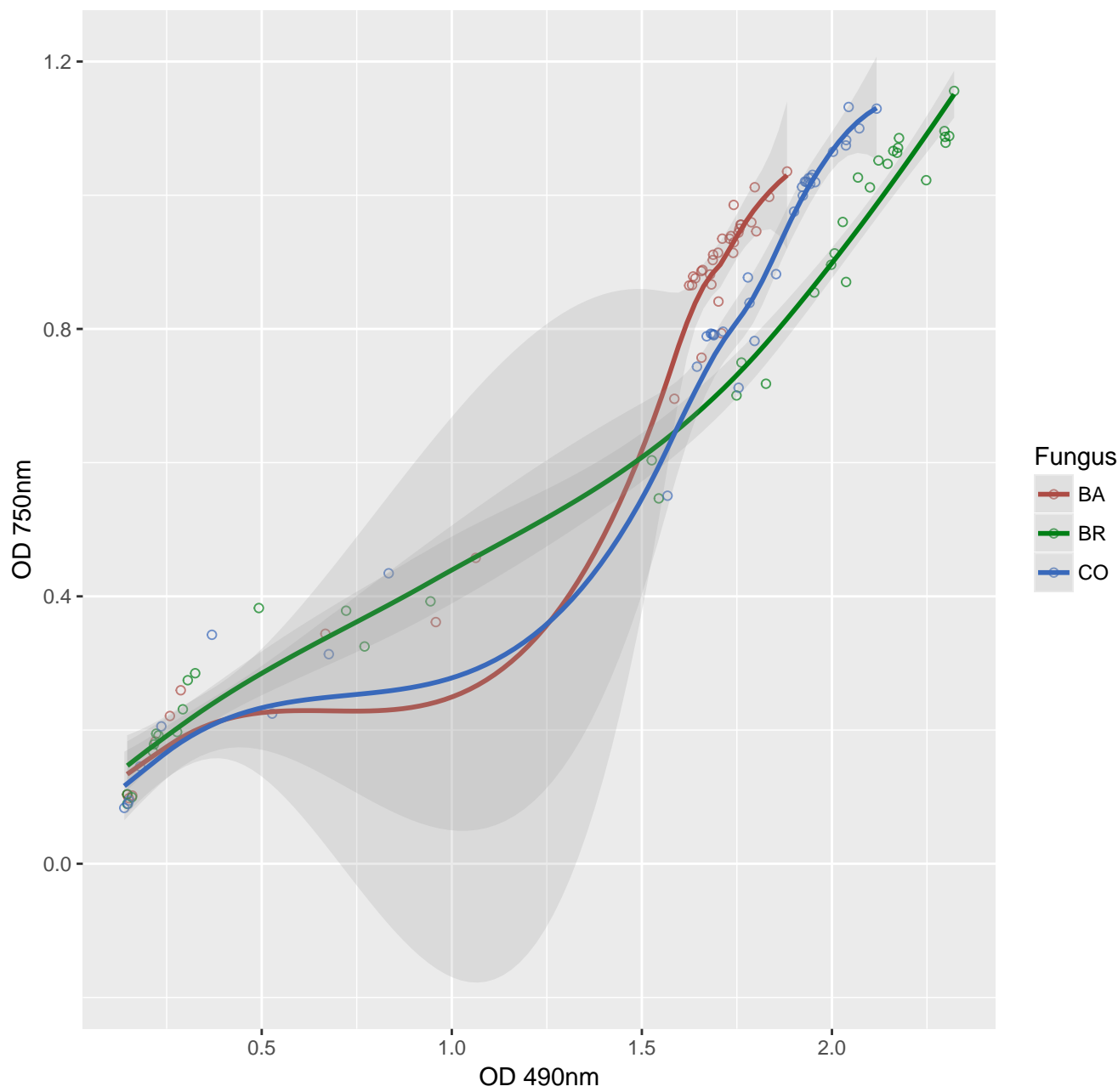
L-Phenylalanine



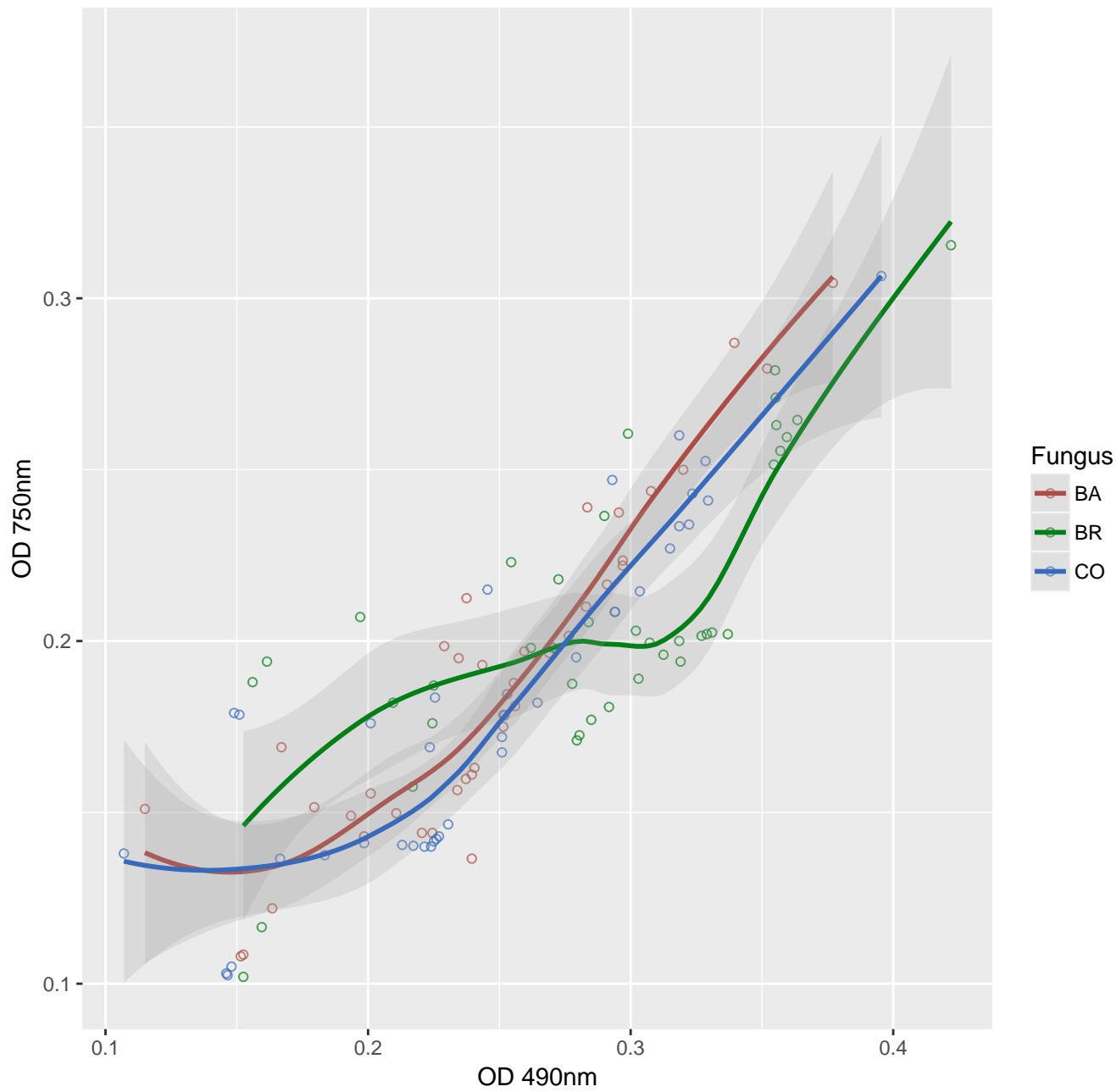
L.Proline



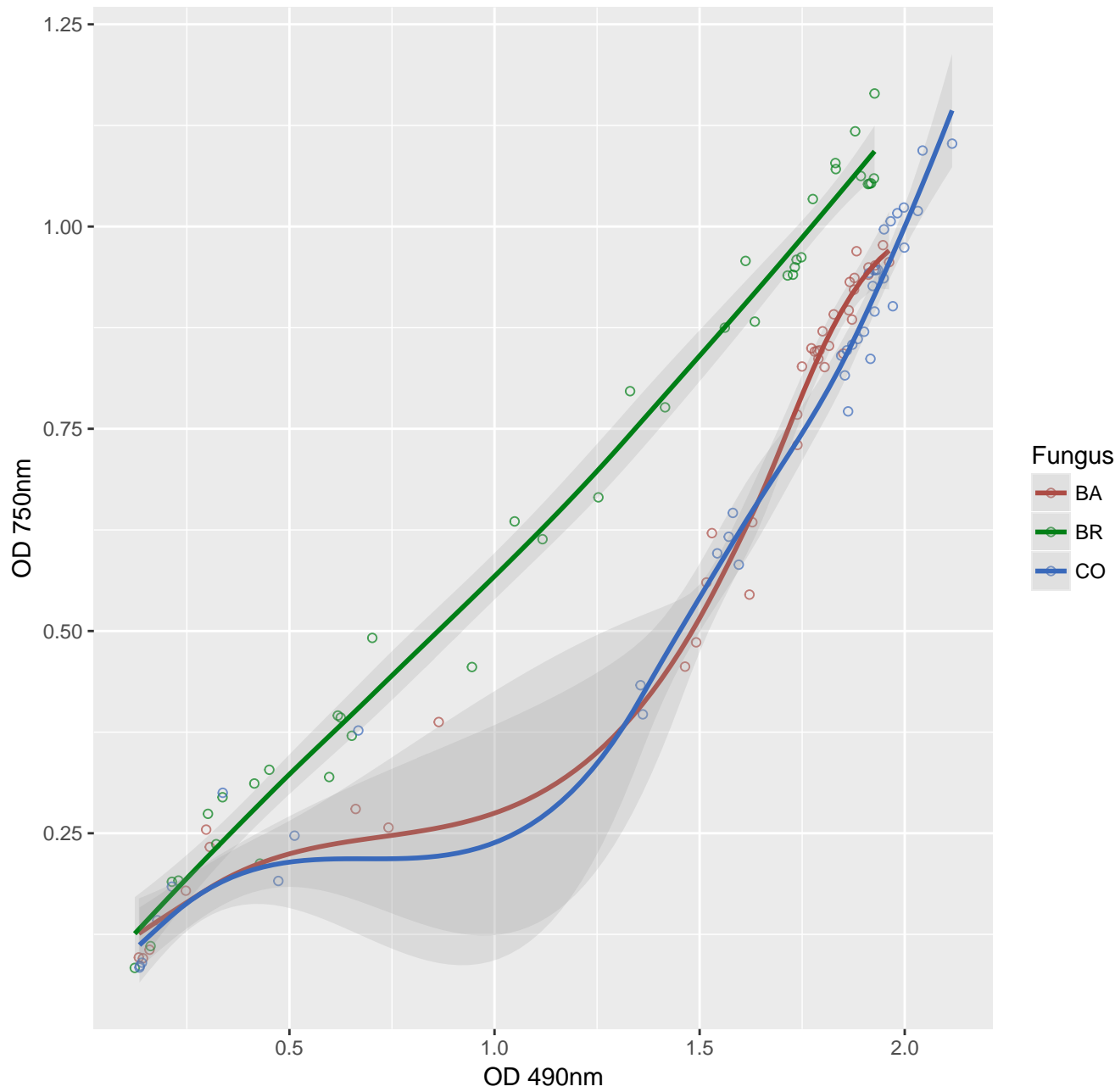
L.PyroglutamicAcid



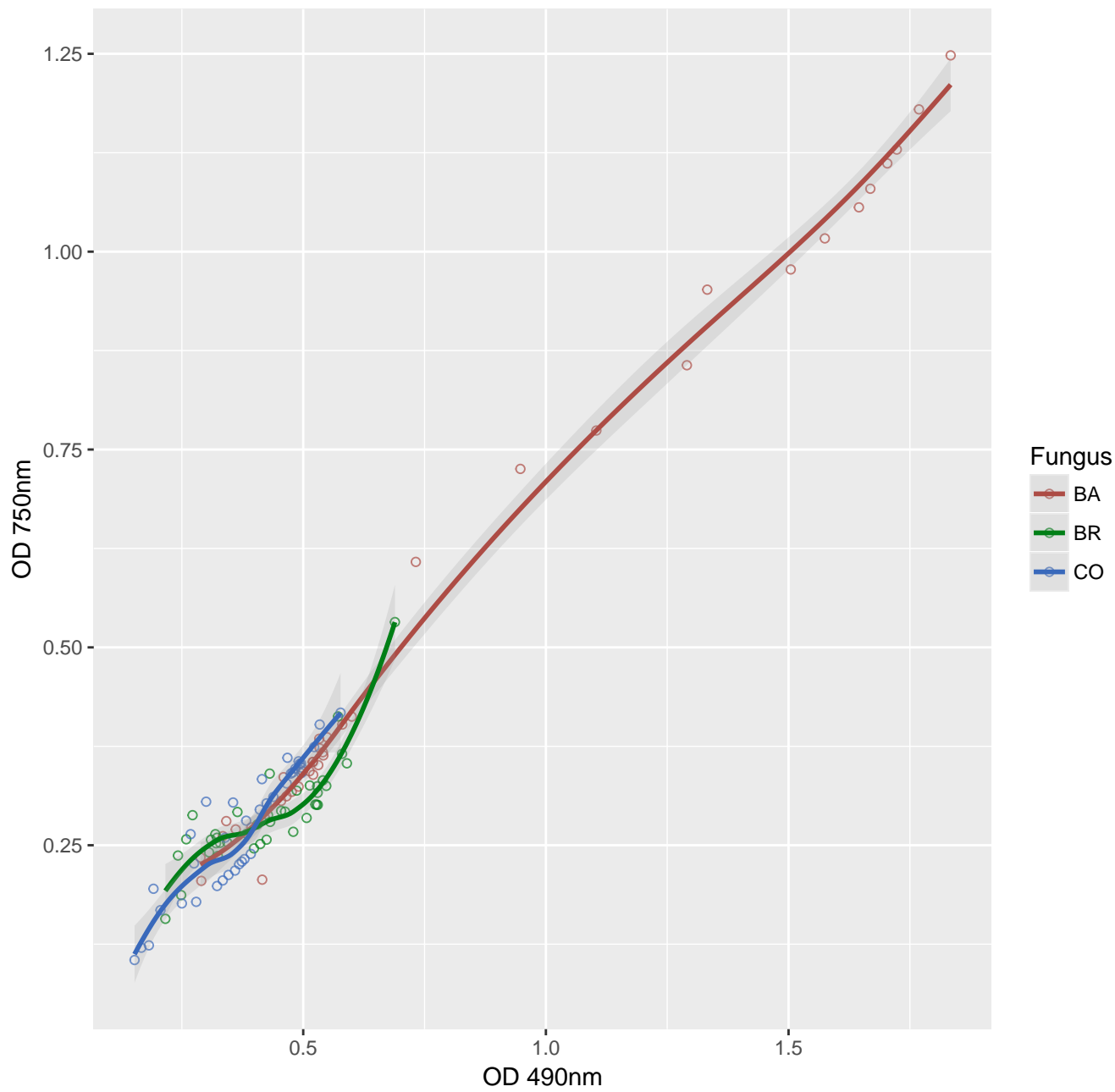
L.Rhamnose



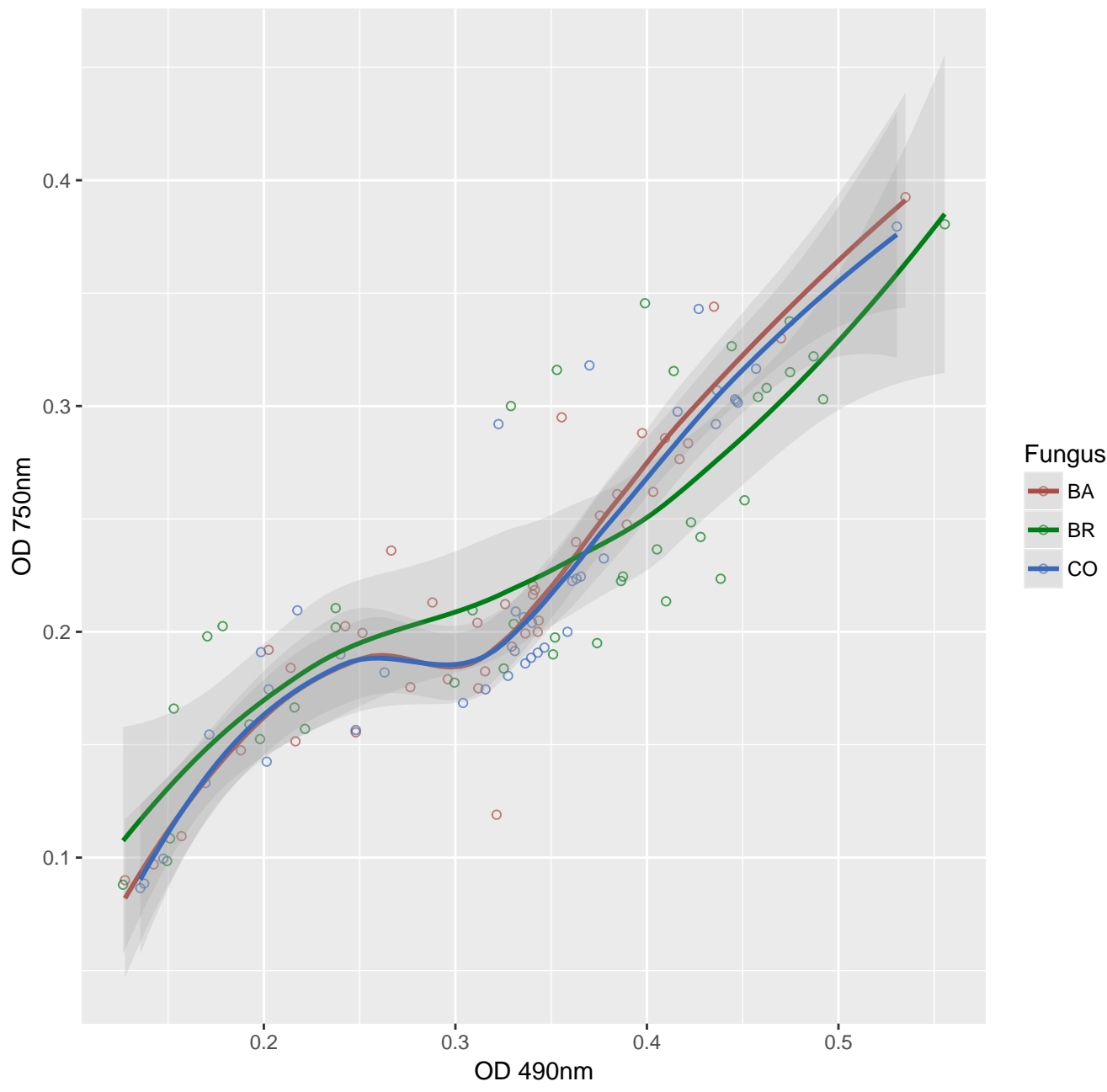
L.Serine



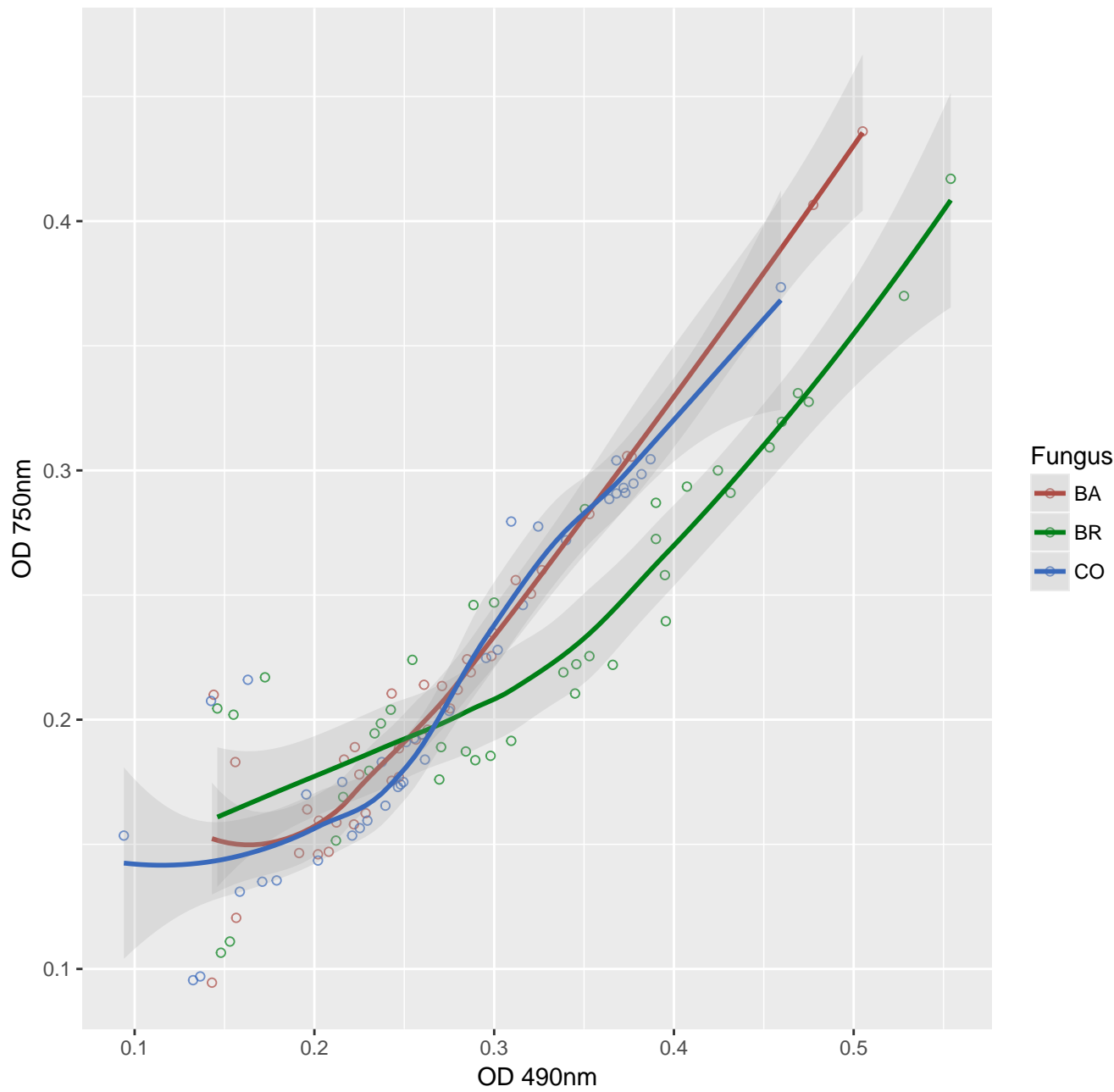
L.Sorbose



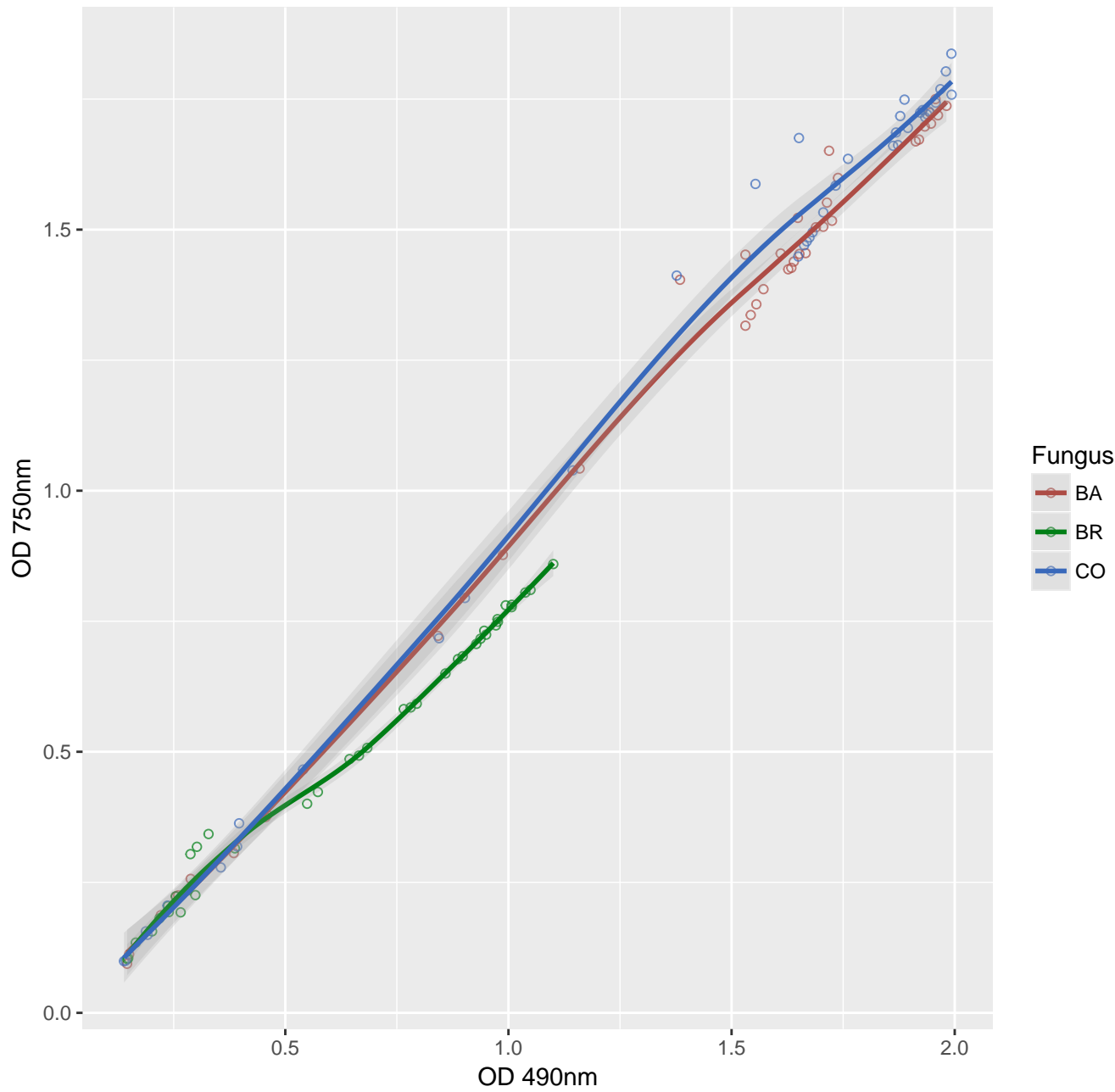
L.Threonine



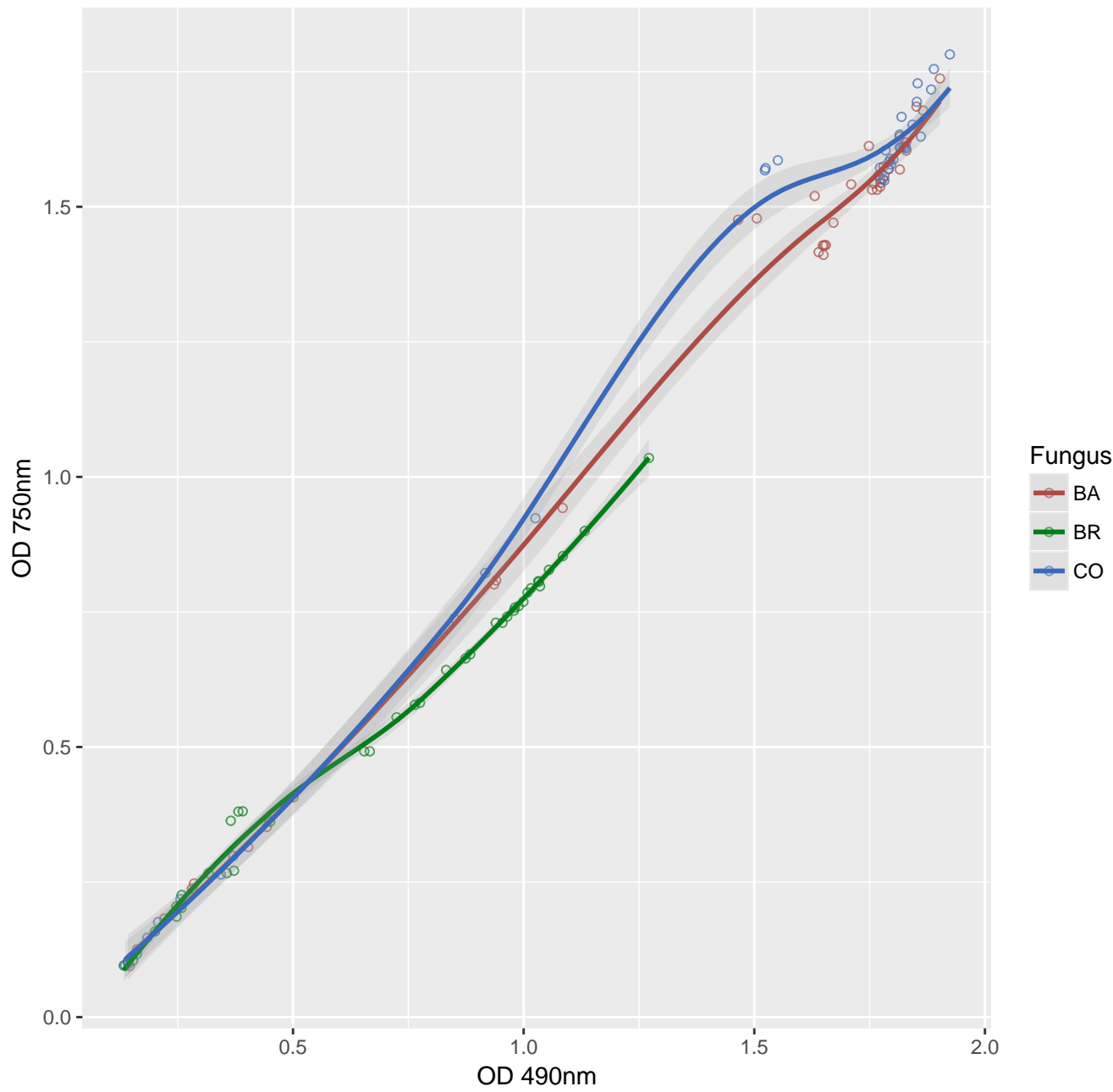
Maltitol



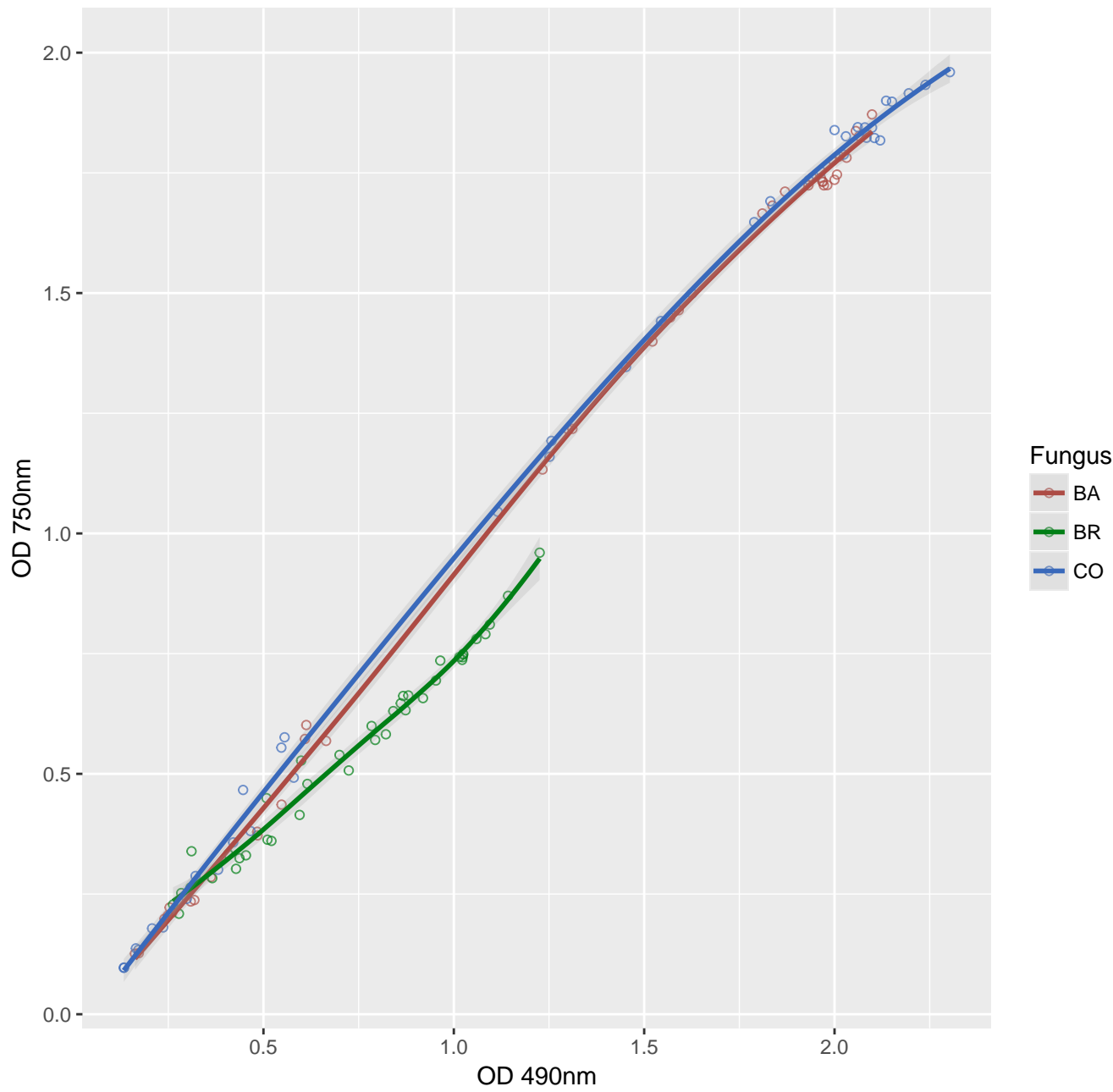
Maltose



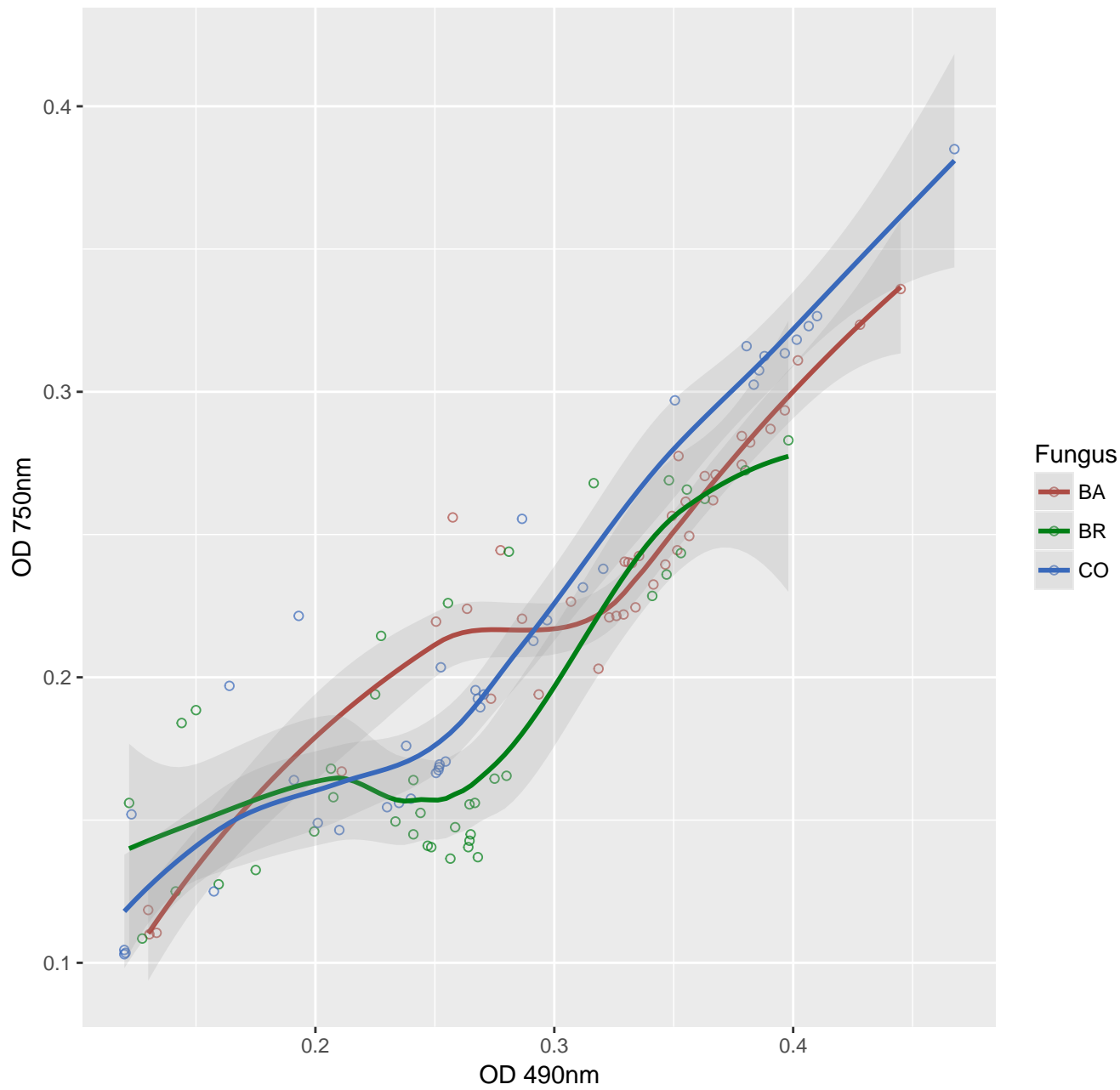
Maltotriose



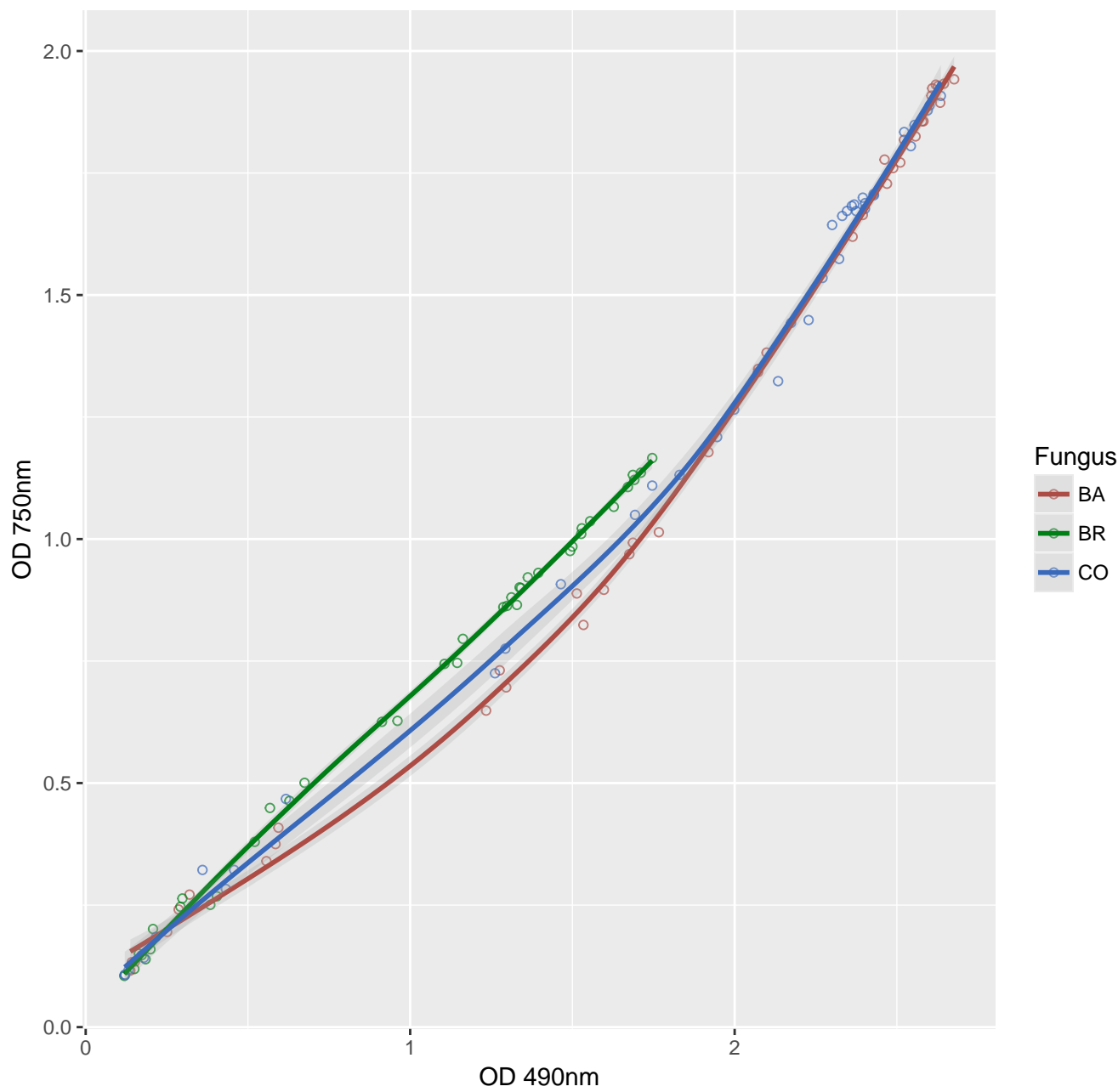
m.Inositol



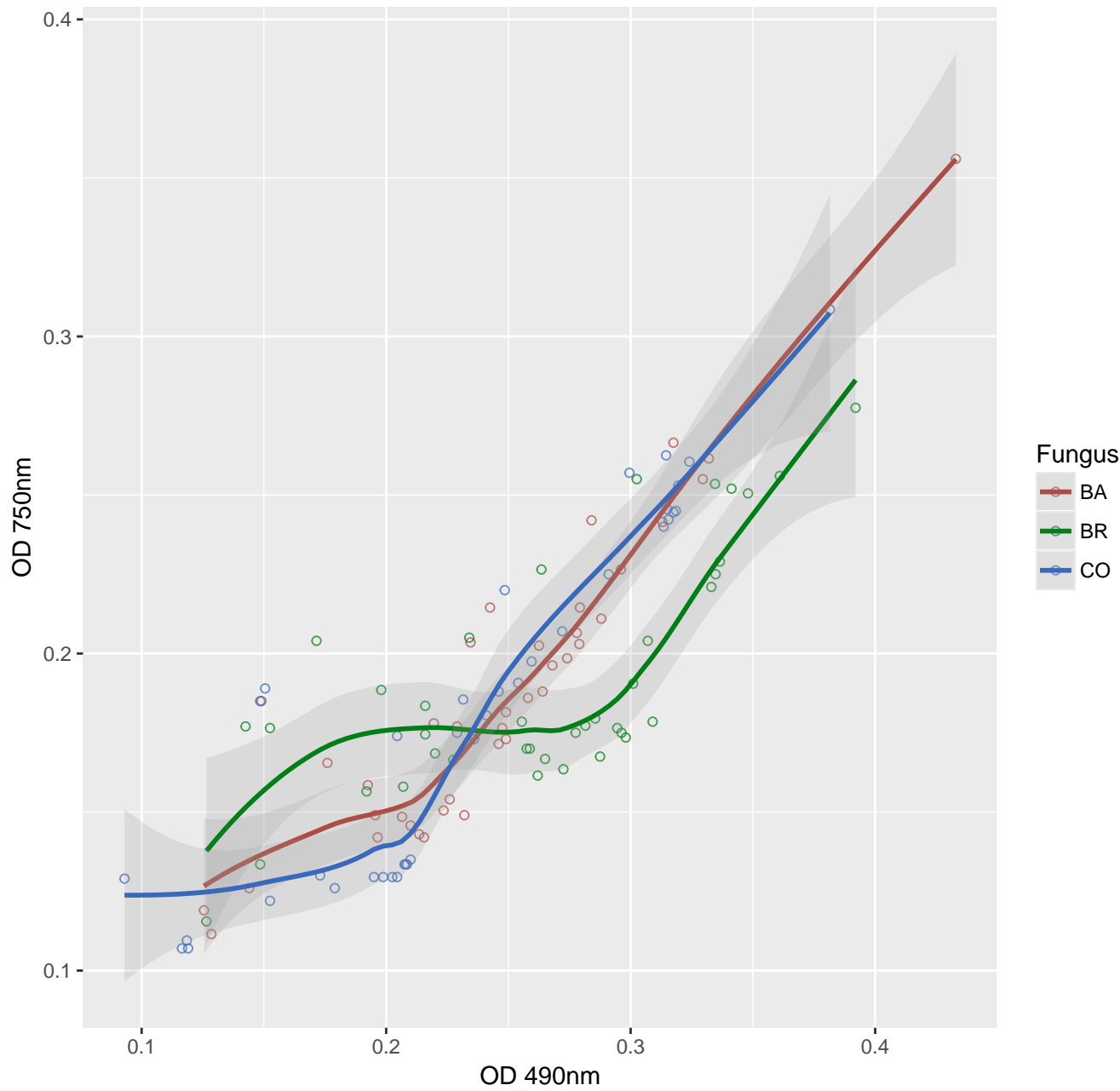
N.Acetyl.D.Galactosamine



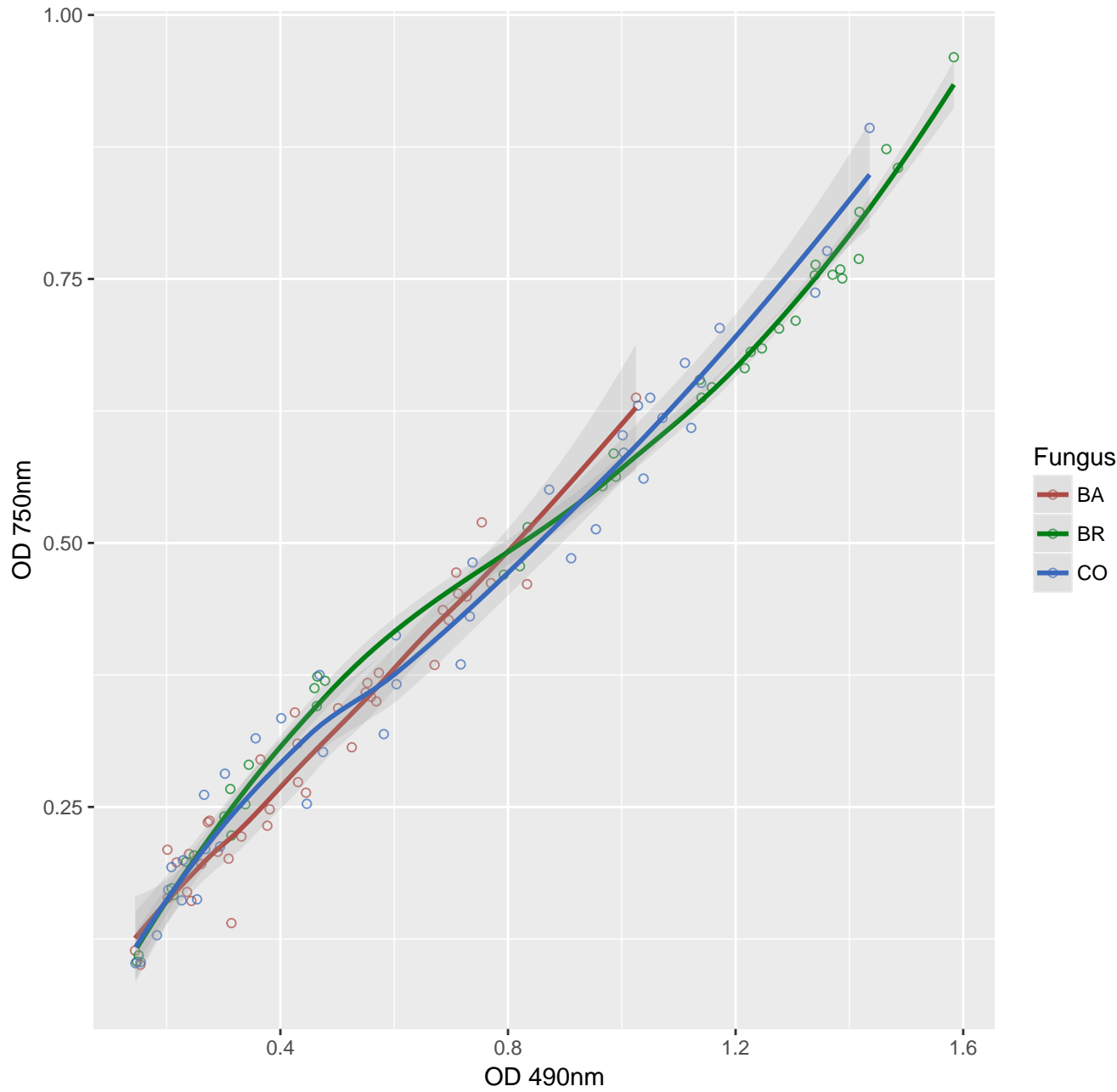
N.Acetyl.D.Glucosamine



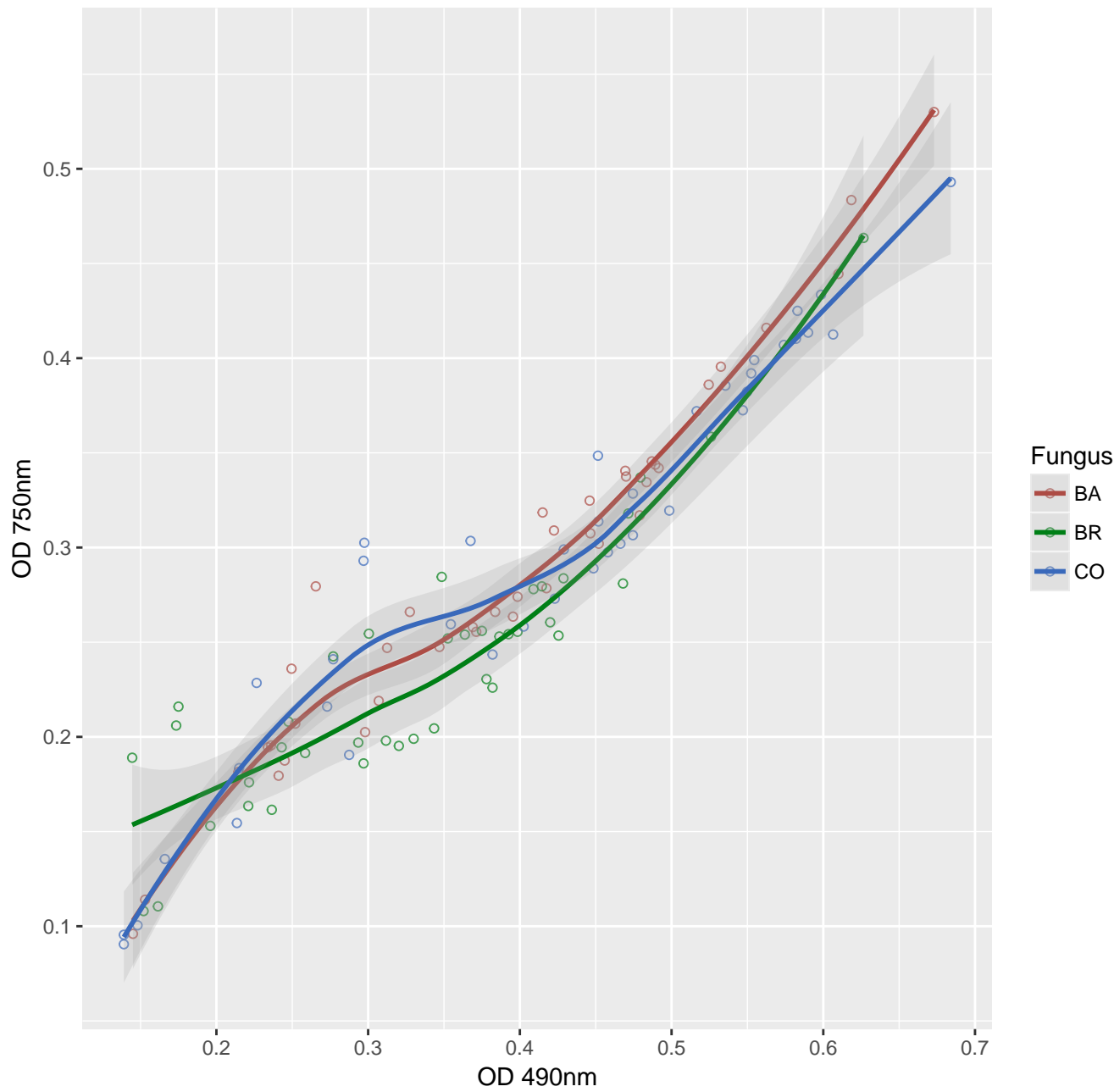
N.Acetyl.D.Mannosamine



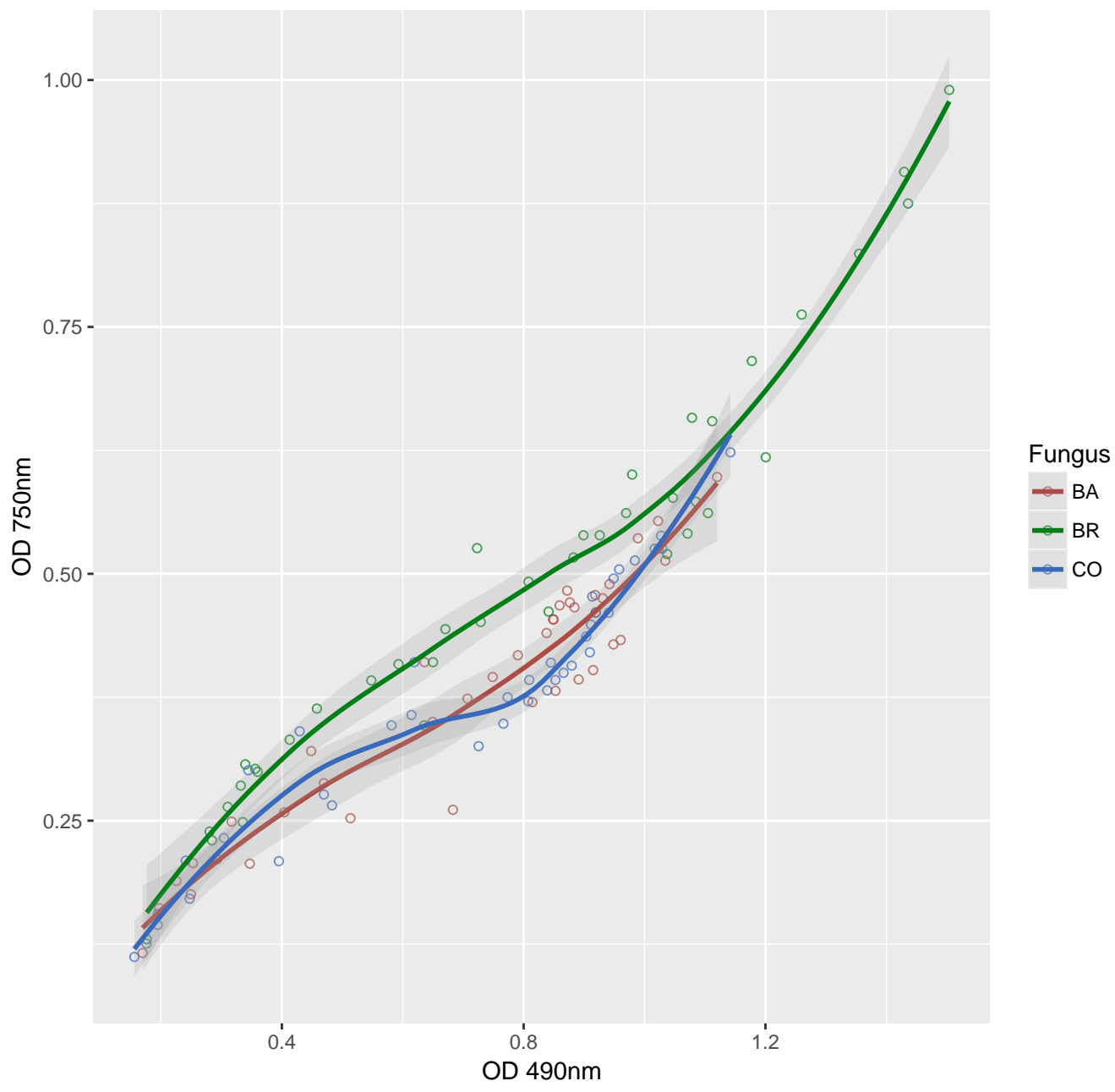
N.Acetyly.L.GlutamicAcid



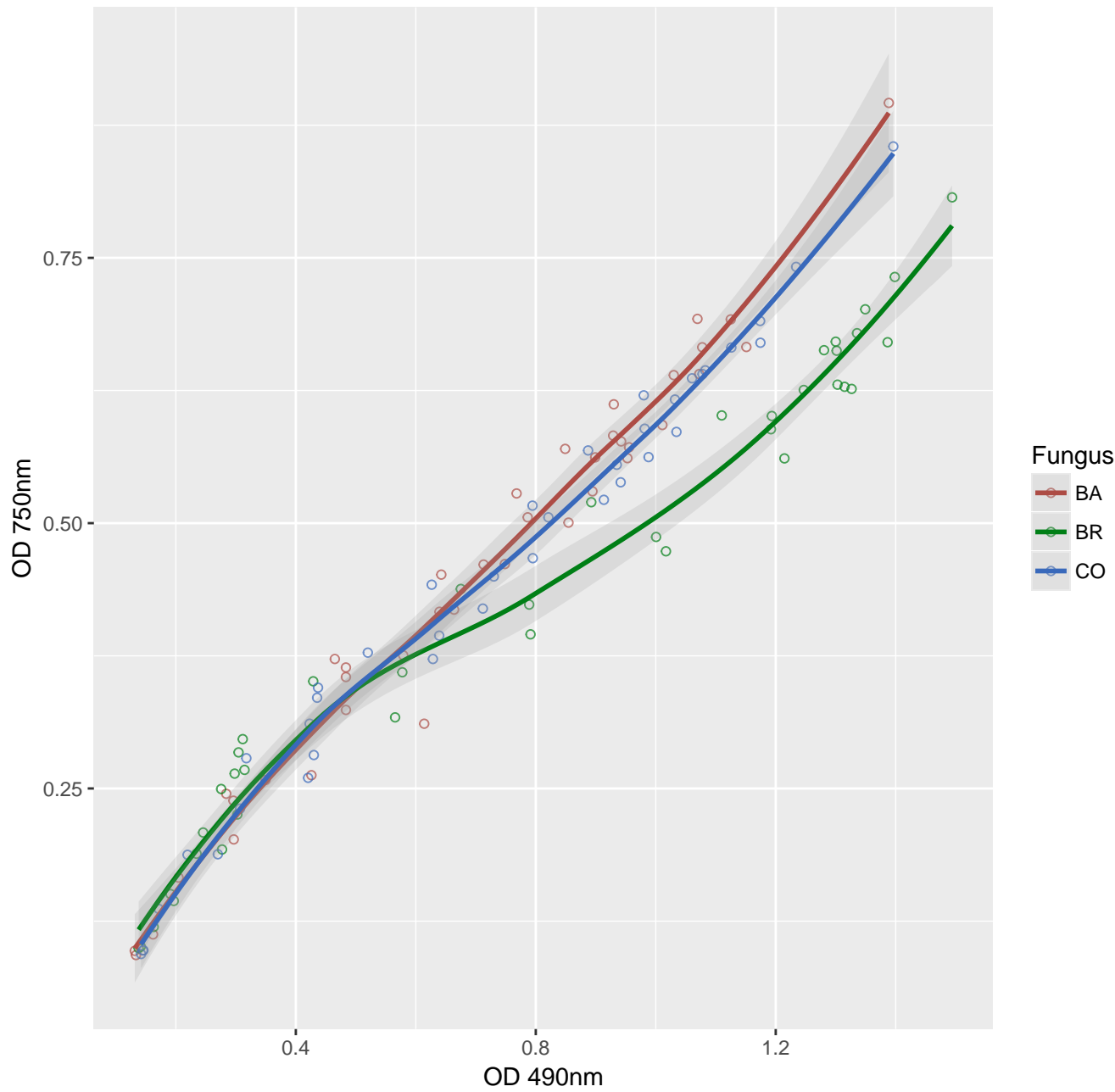
Palatinose



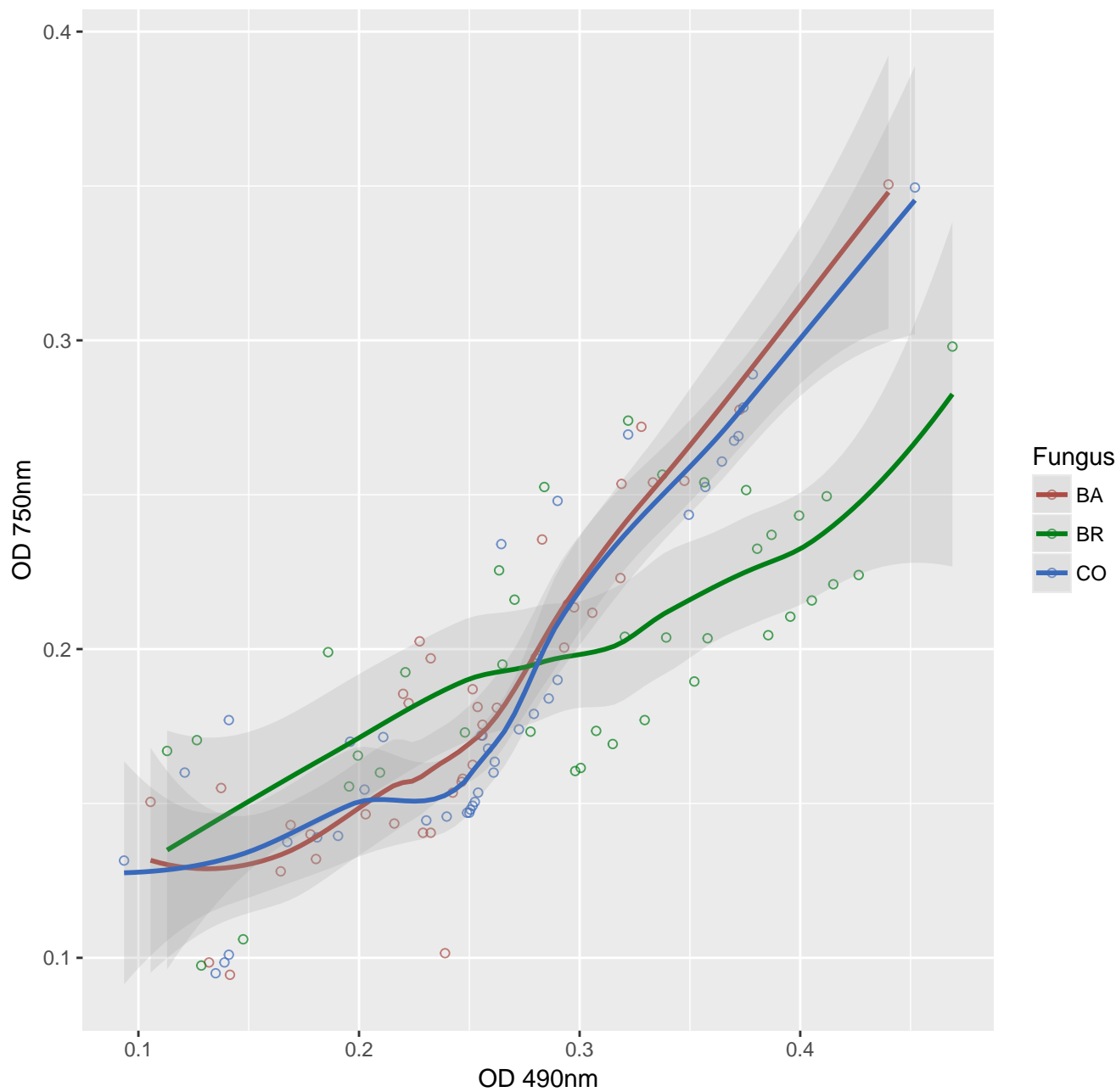
P.Hydroxyphenyl.aceticAcid



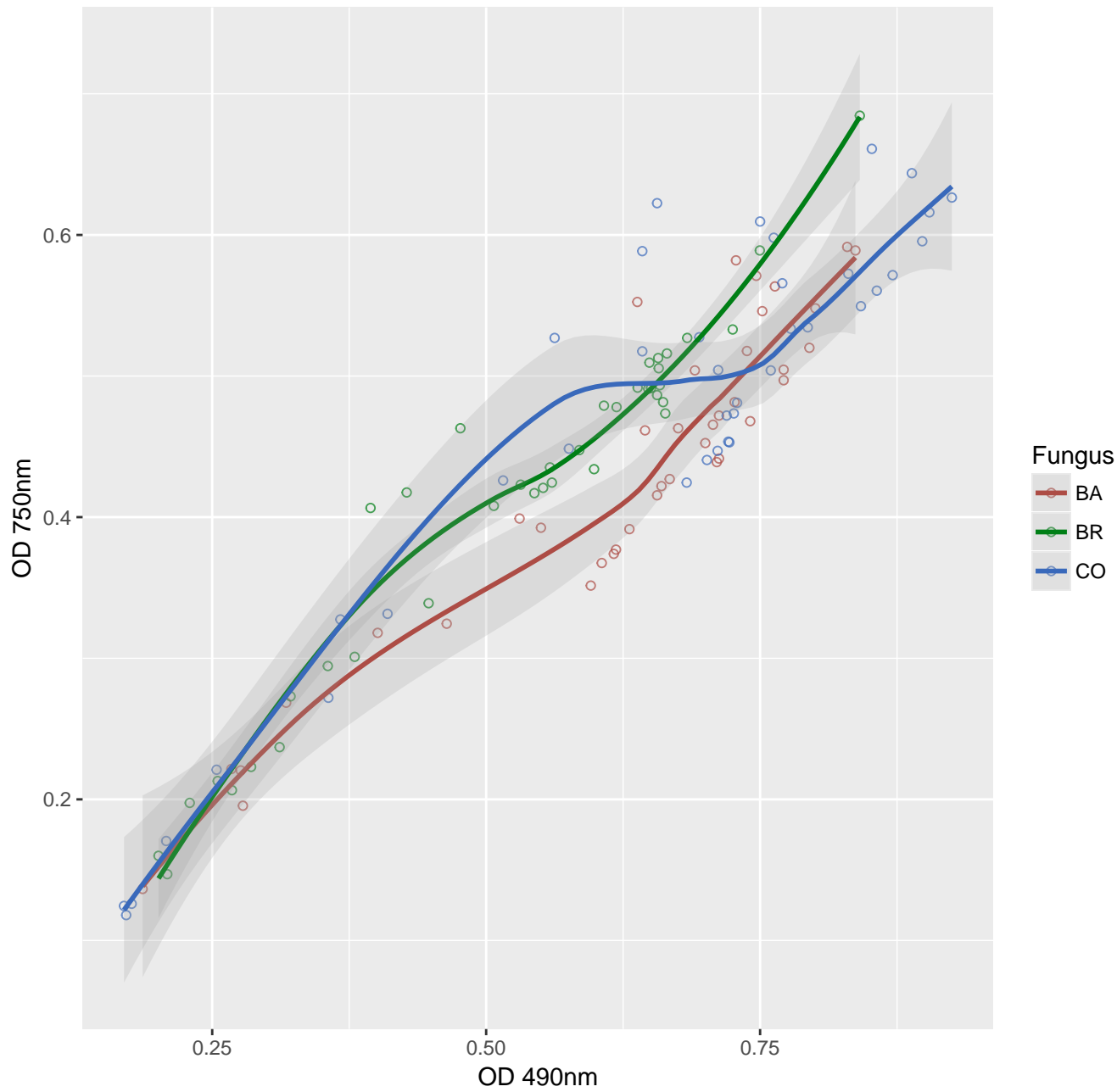
Putrescine



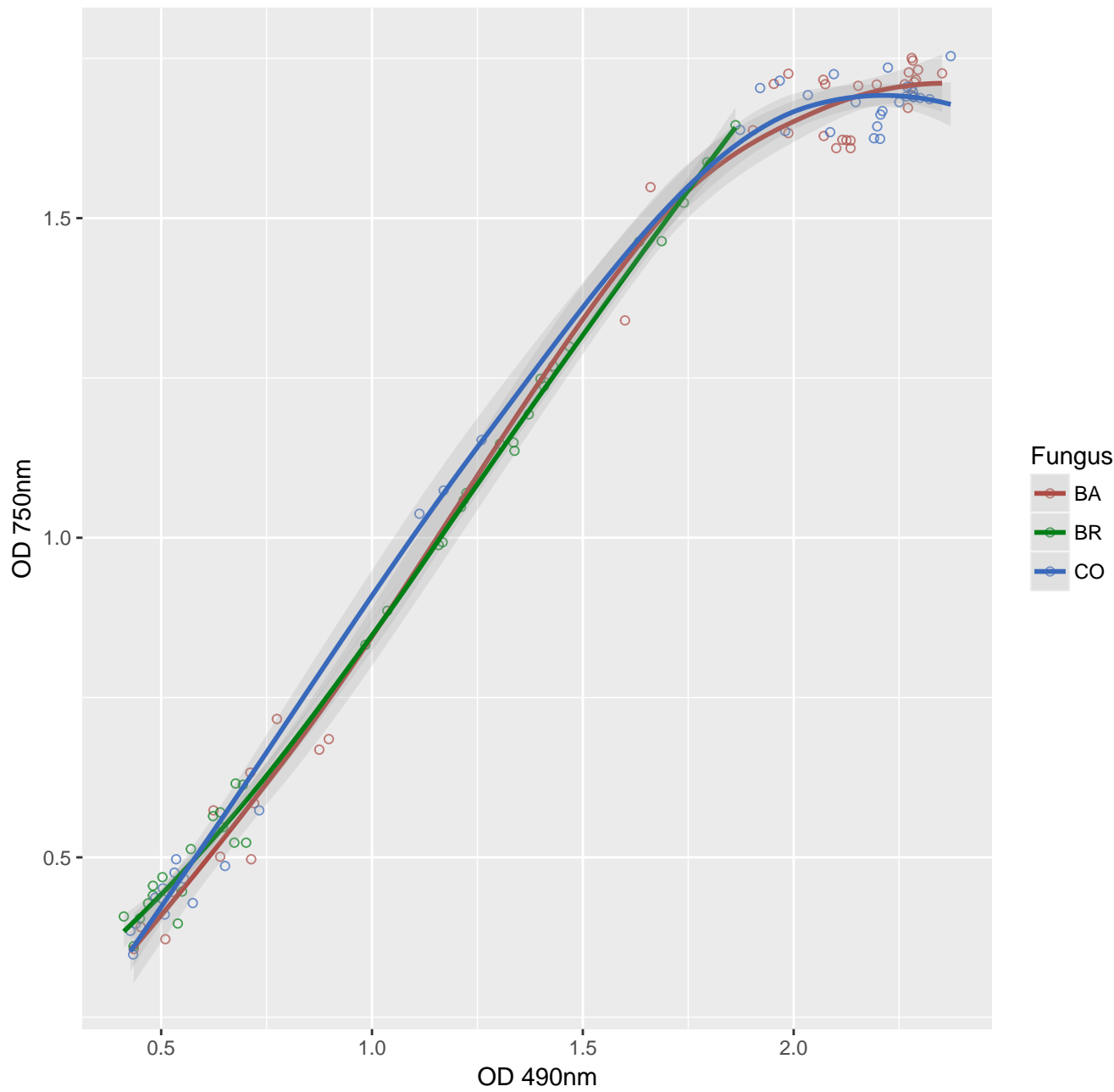
QuinicAcid



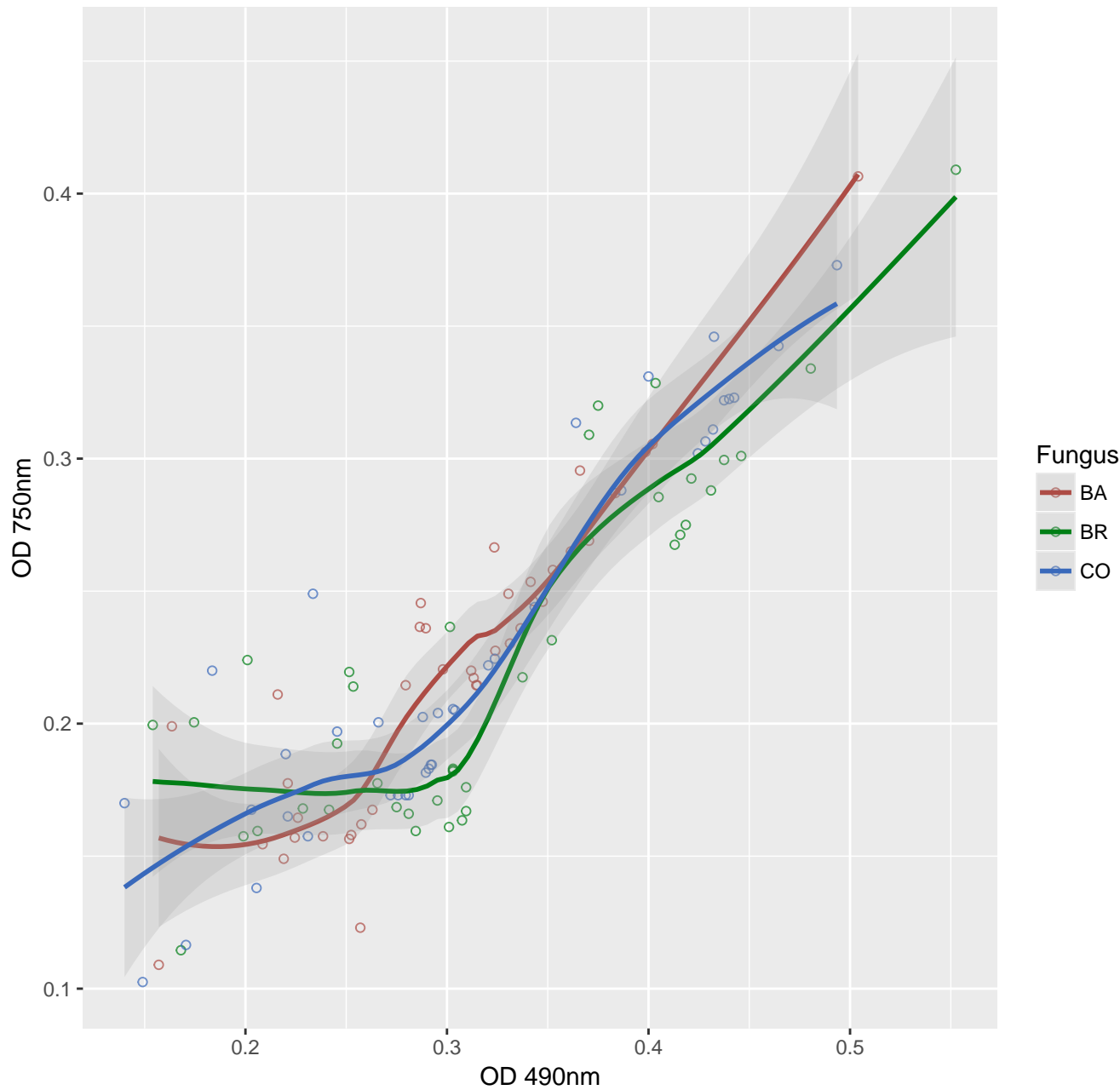
Salicin



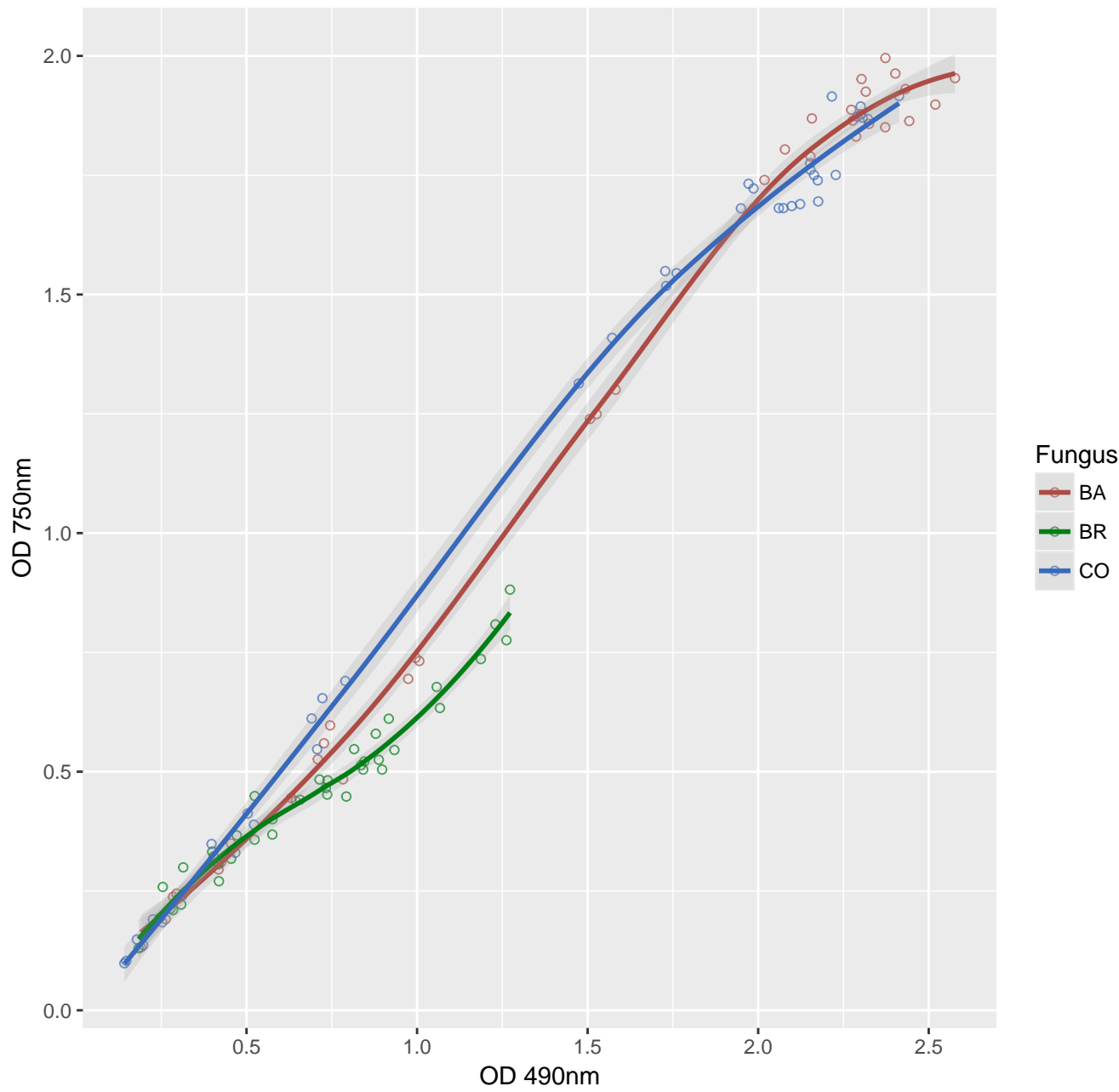
SebacicAcid



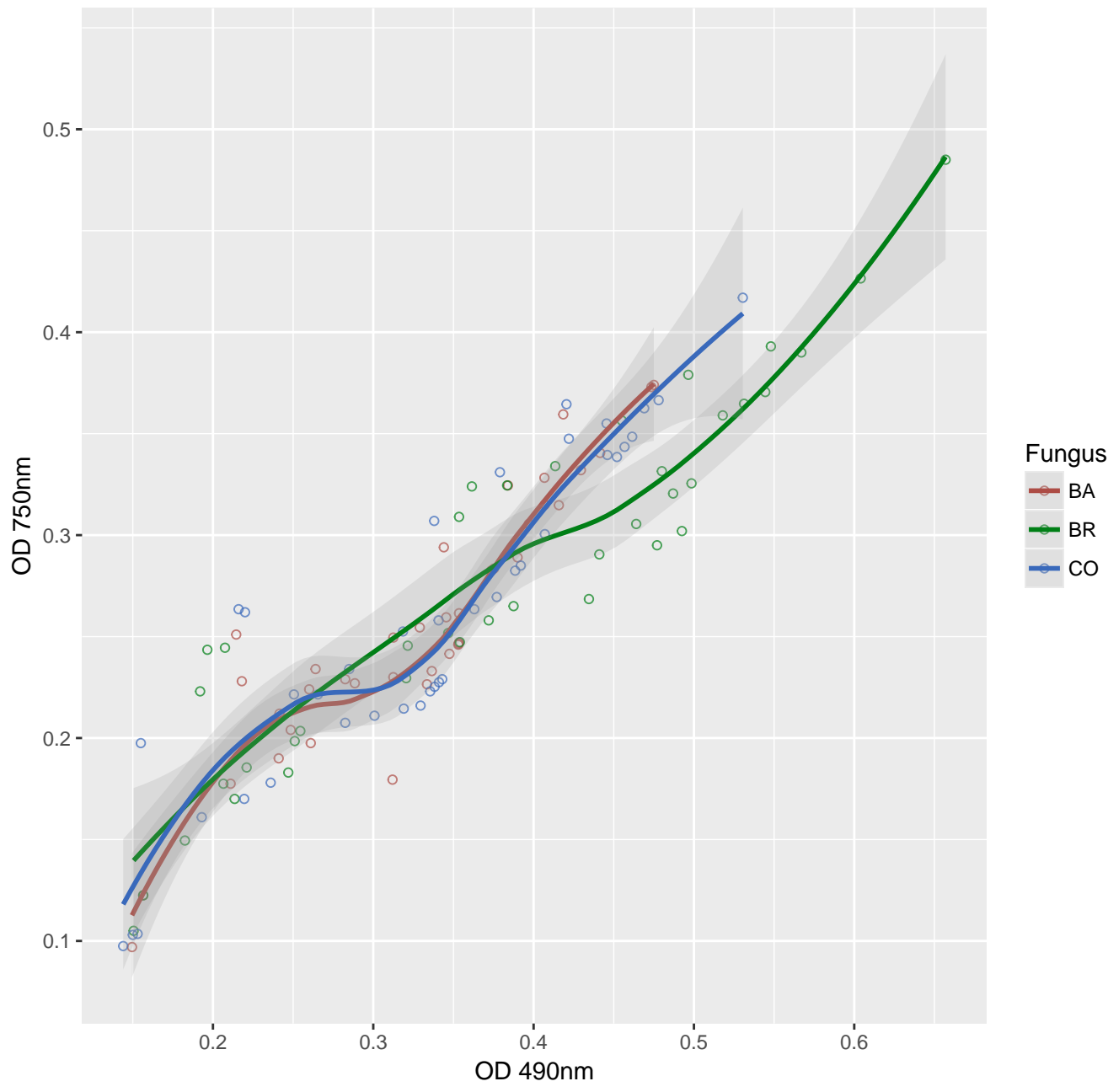
Sedoheptulosan



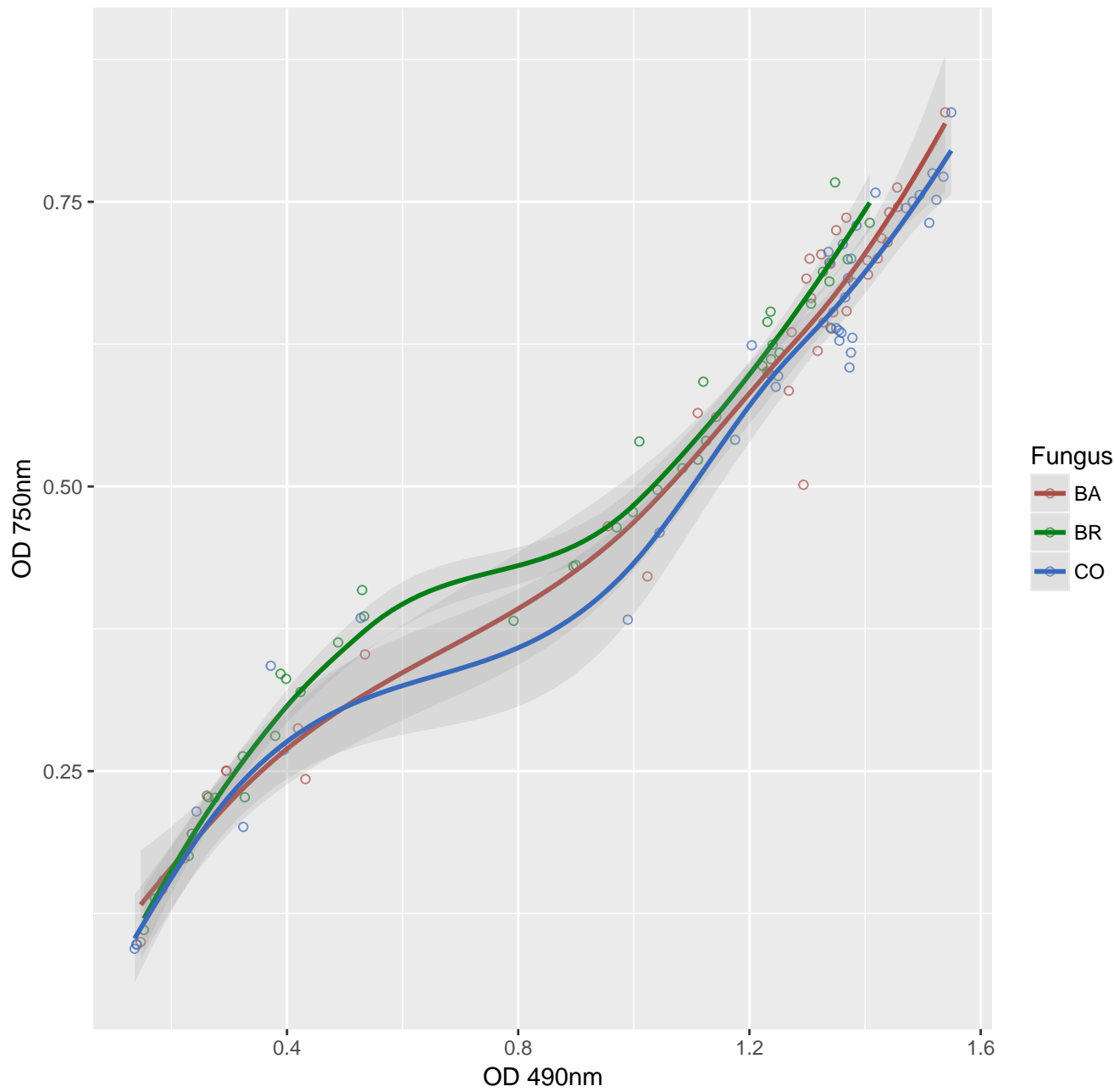
Stachynose



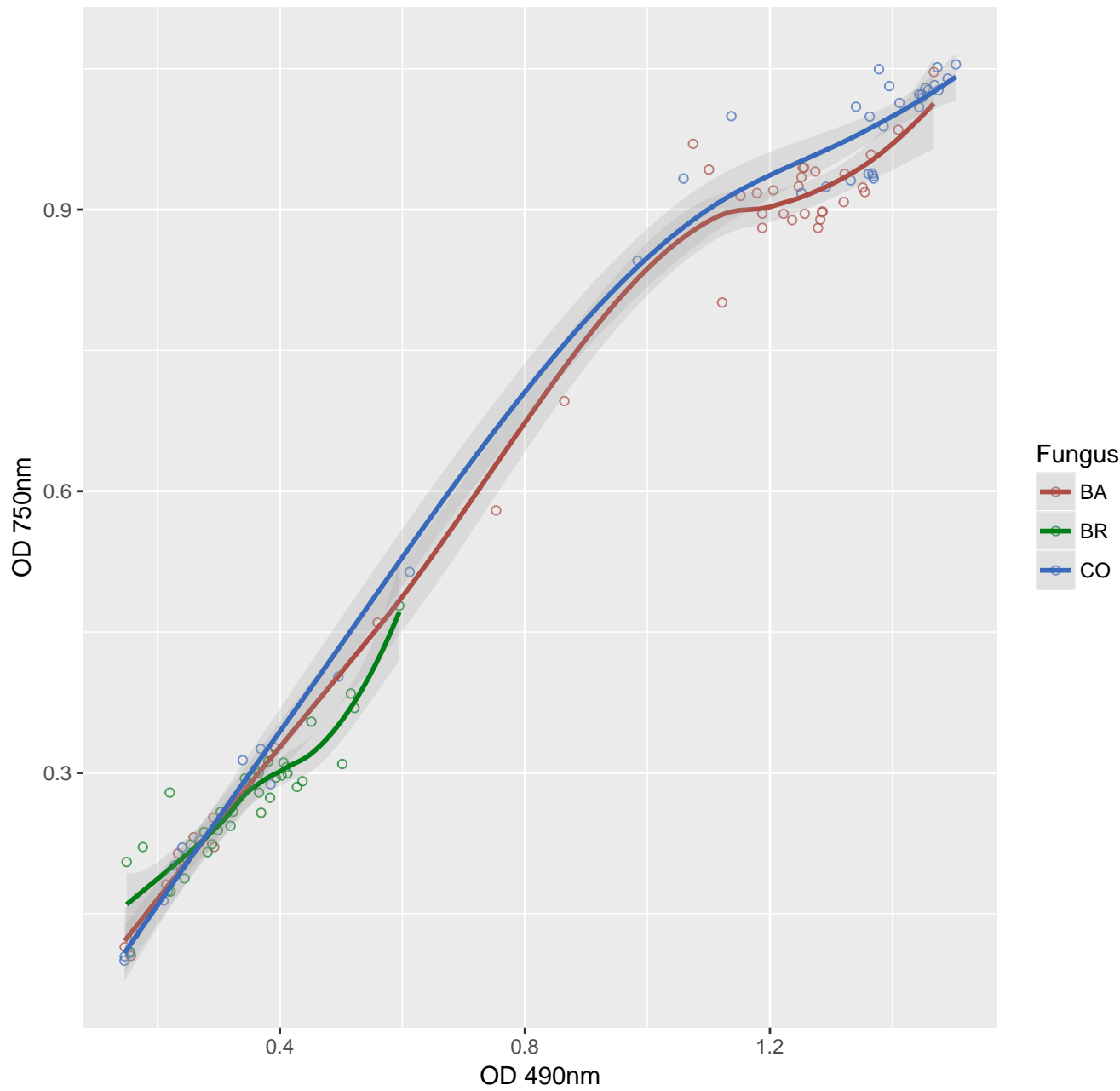
SuccinamicAcid



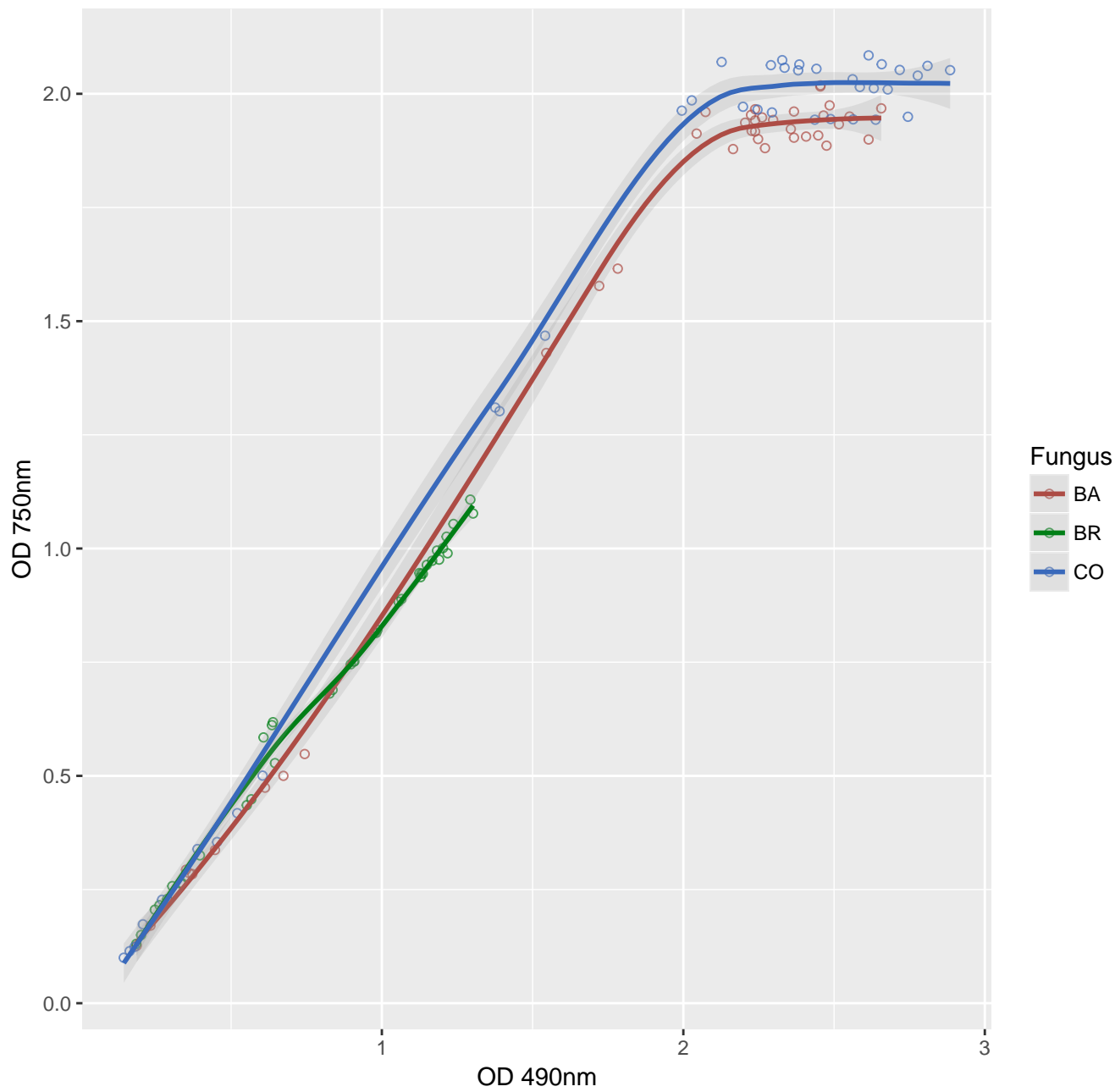
SuccinicAcid



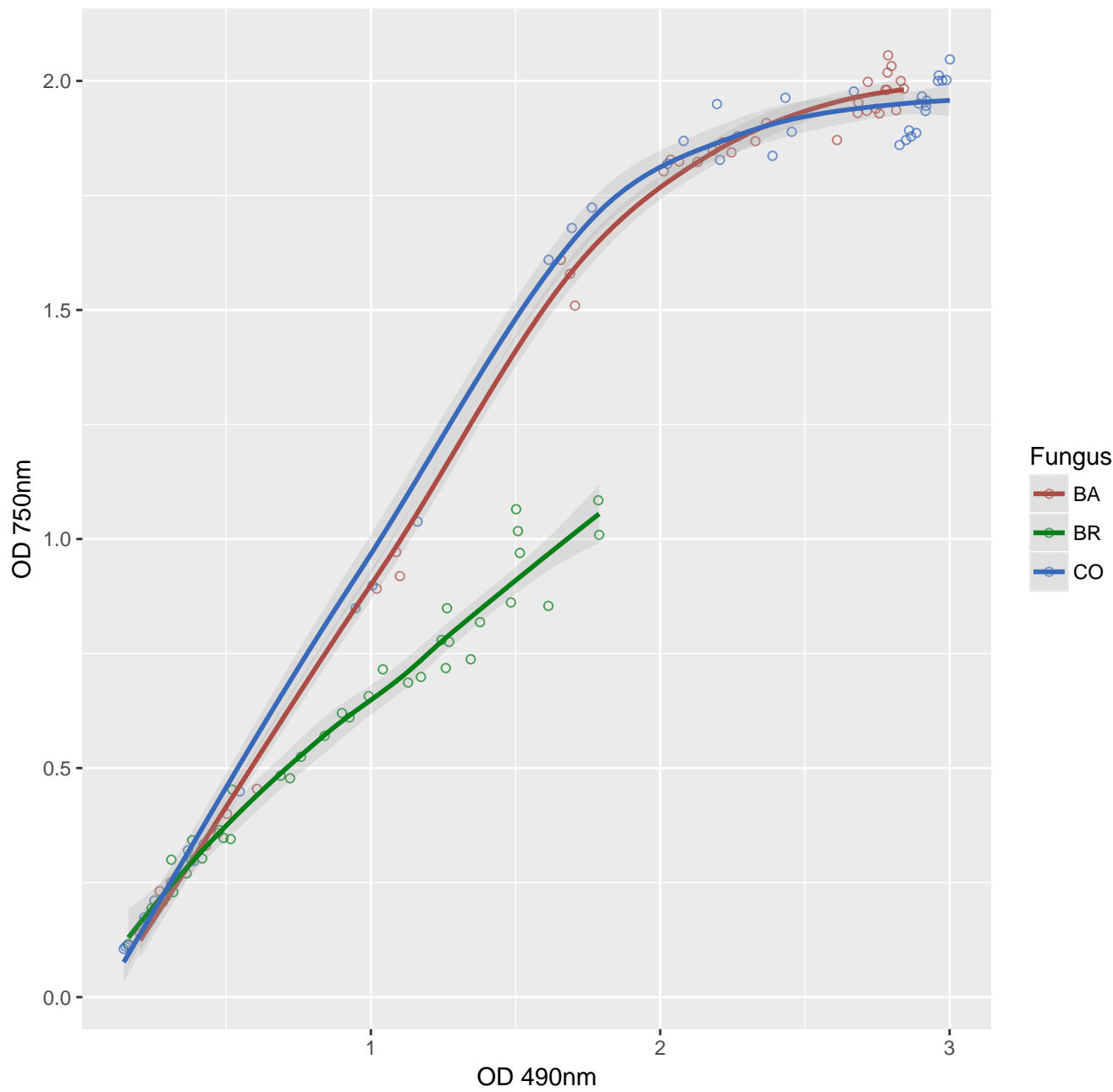
SuccinicAcidMono.MethylEster



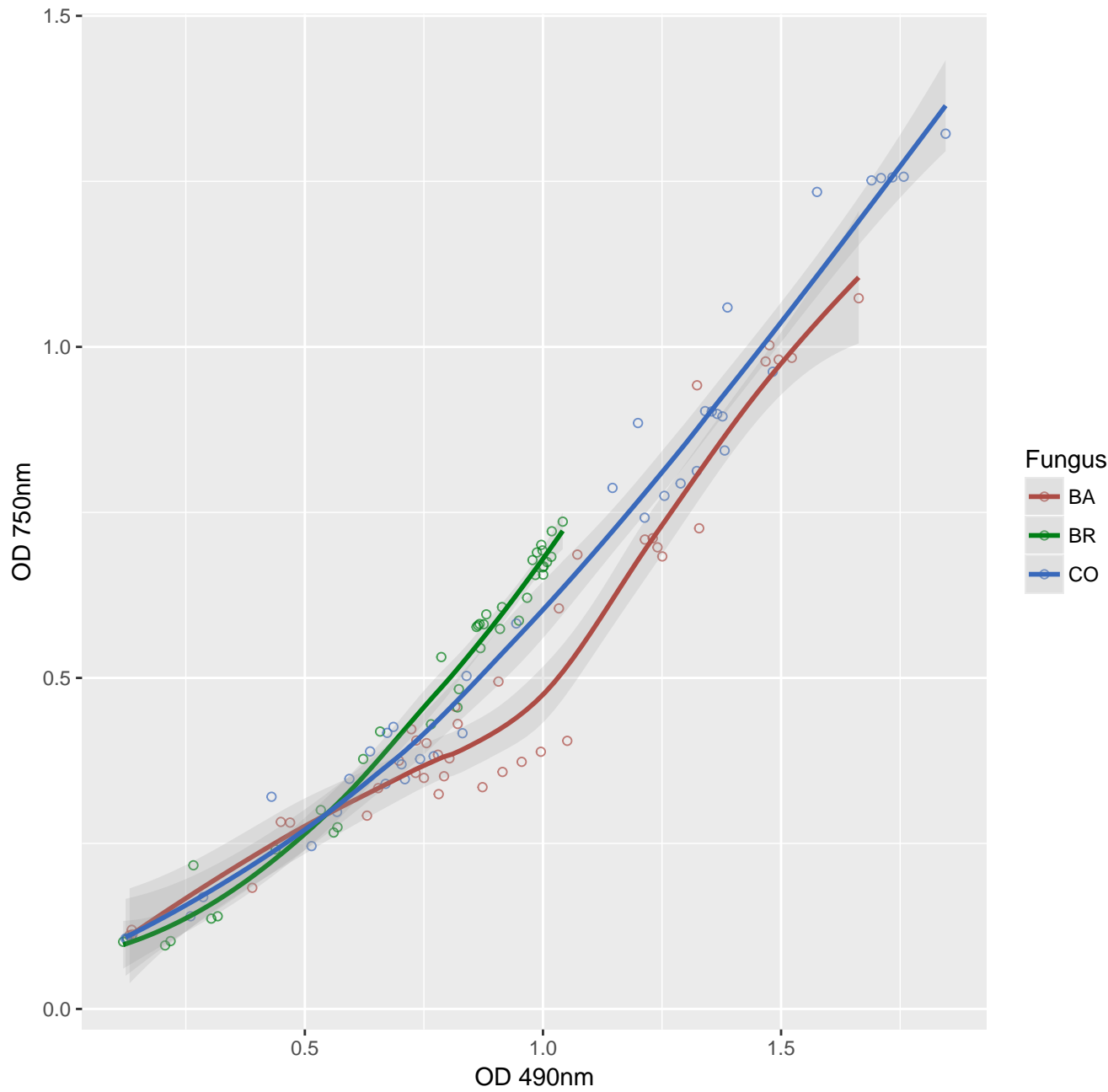
Sucrose



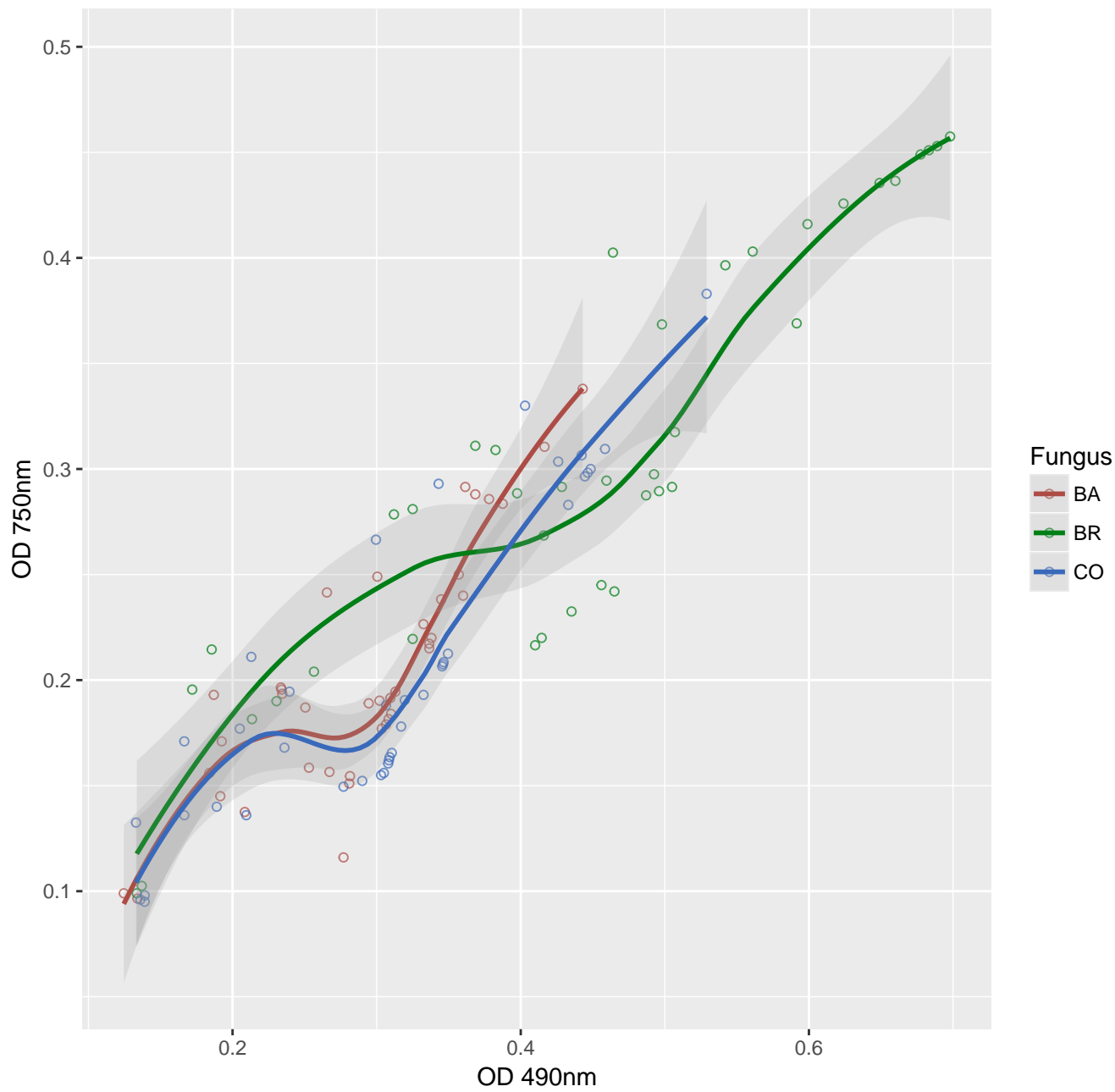
Turanose



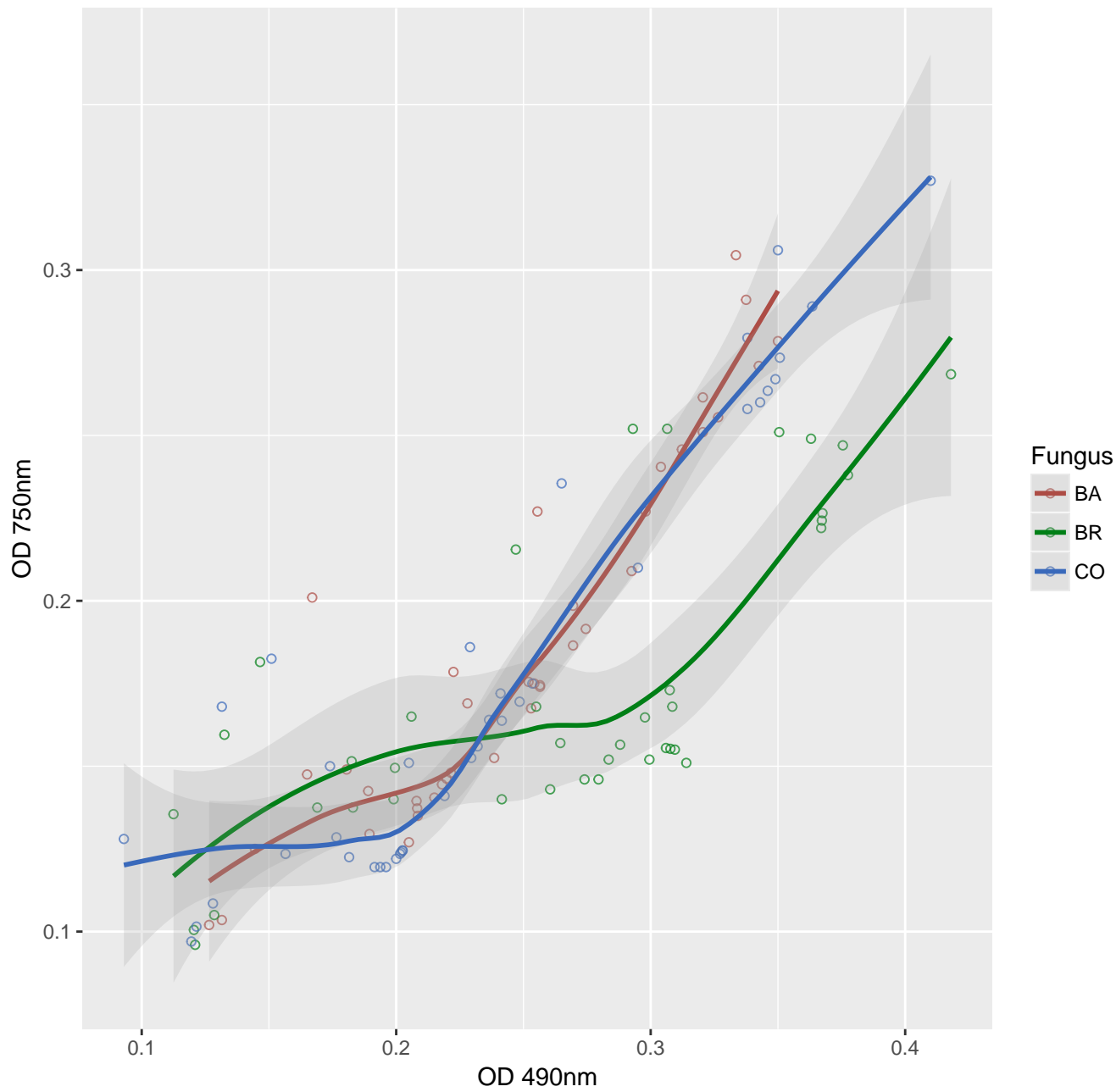
Tween80



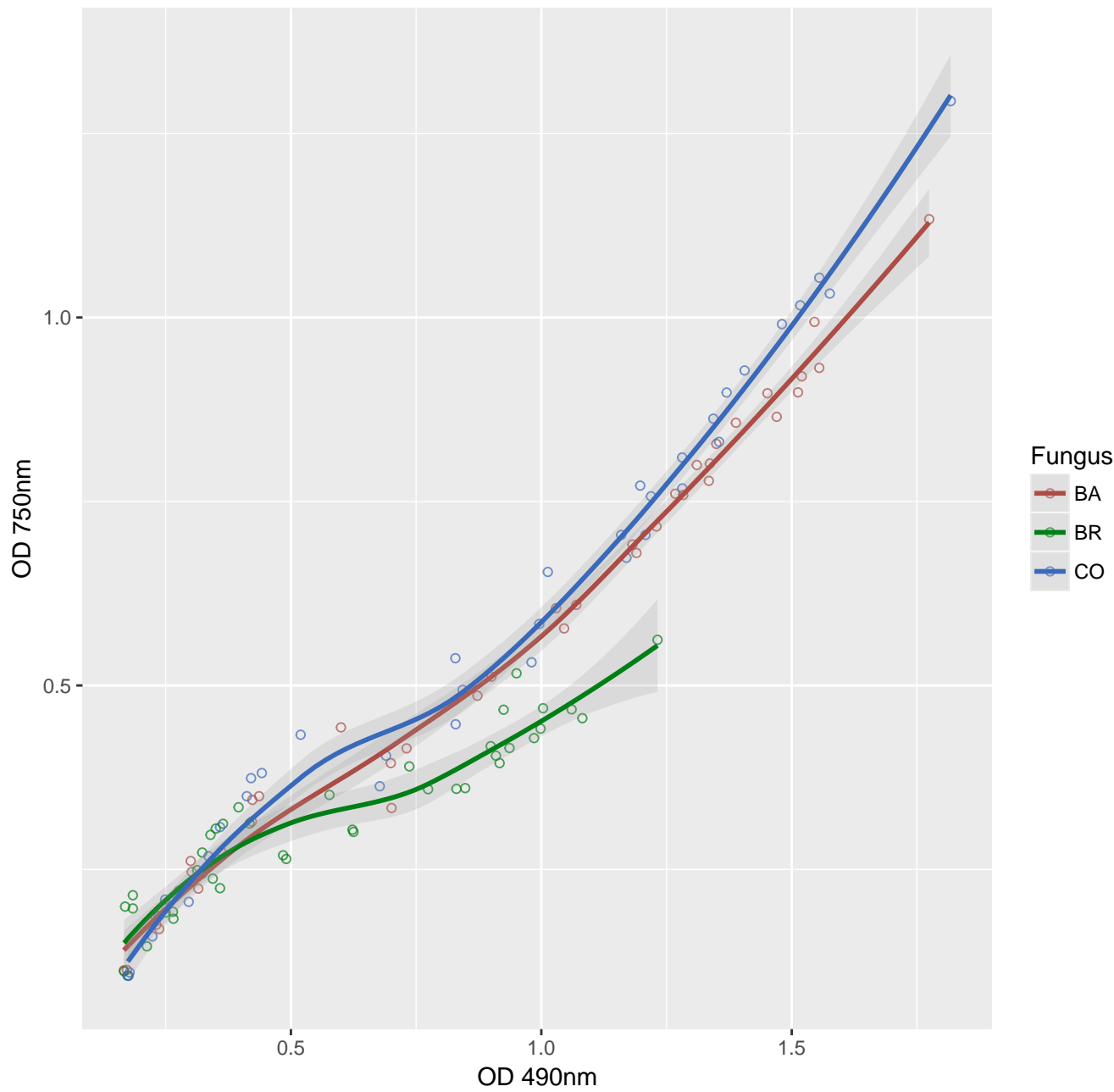
Uridine



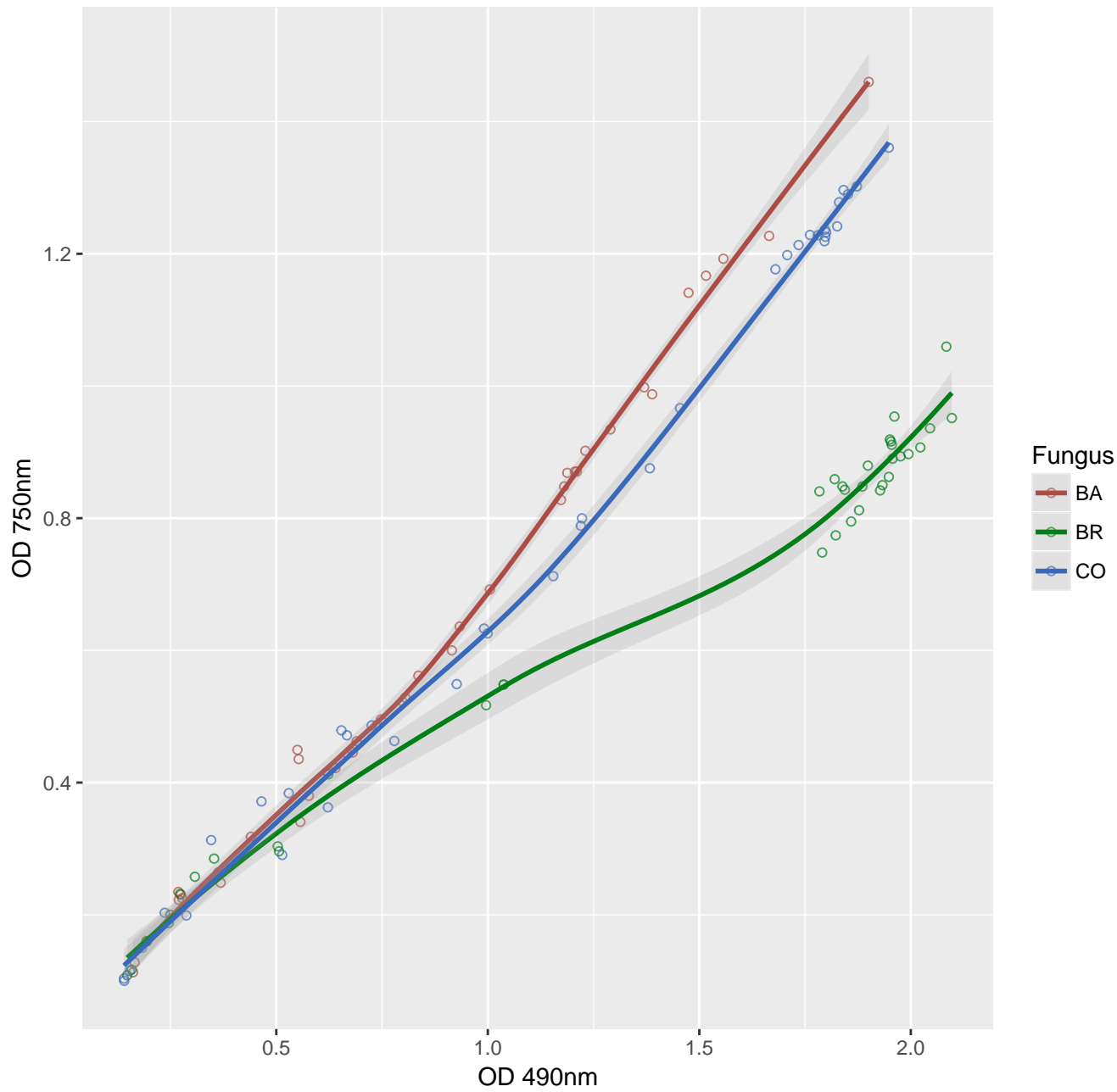
Water



X2.AminoEthanol



X2.Keto.D.GluconicAcid



Xylitol

