

Similarity-Based Multimodal Regression Supplementary Materials

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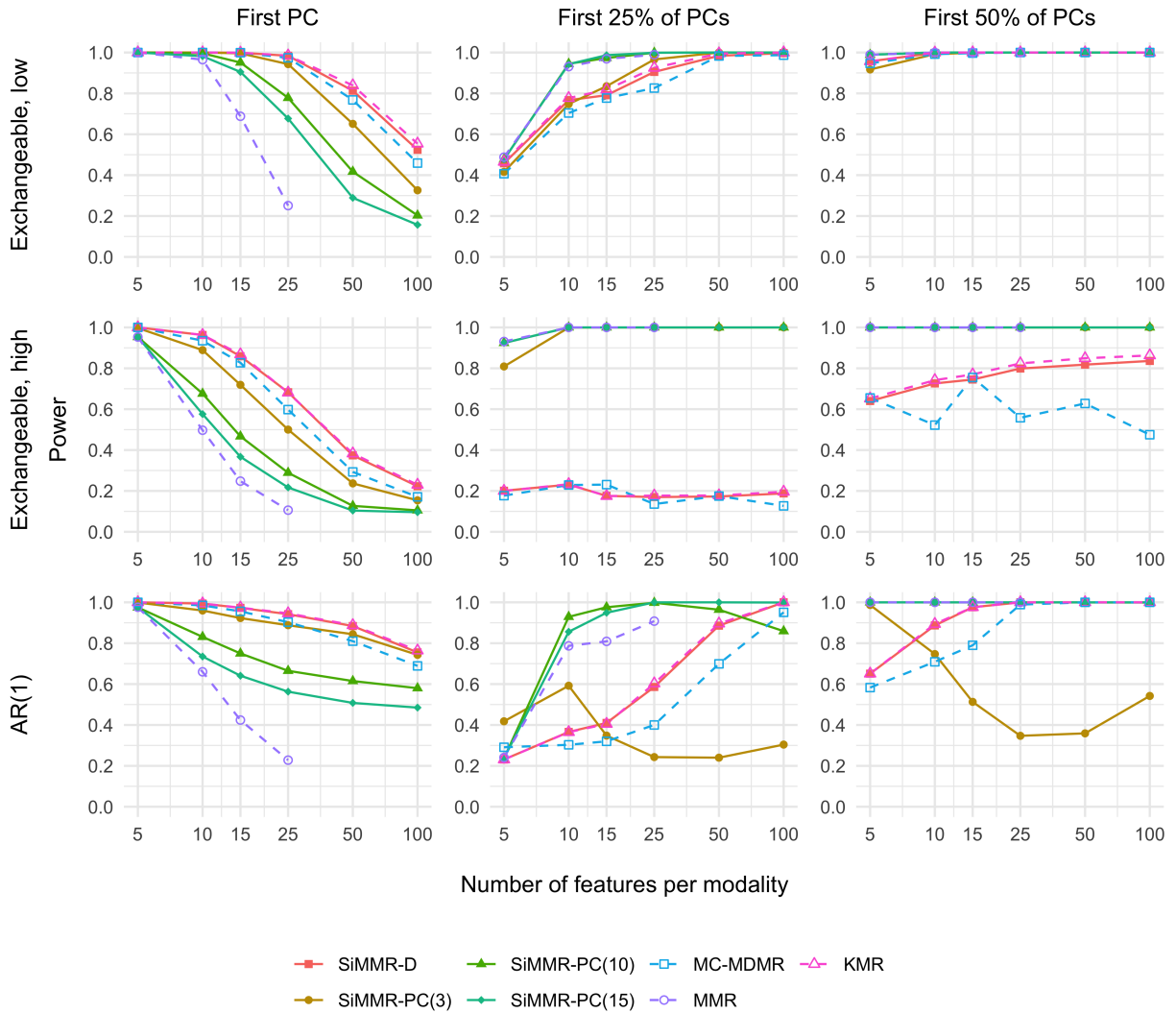
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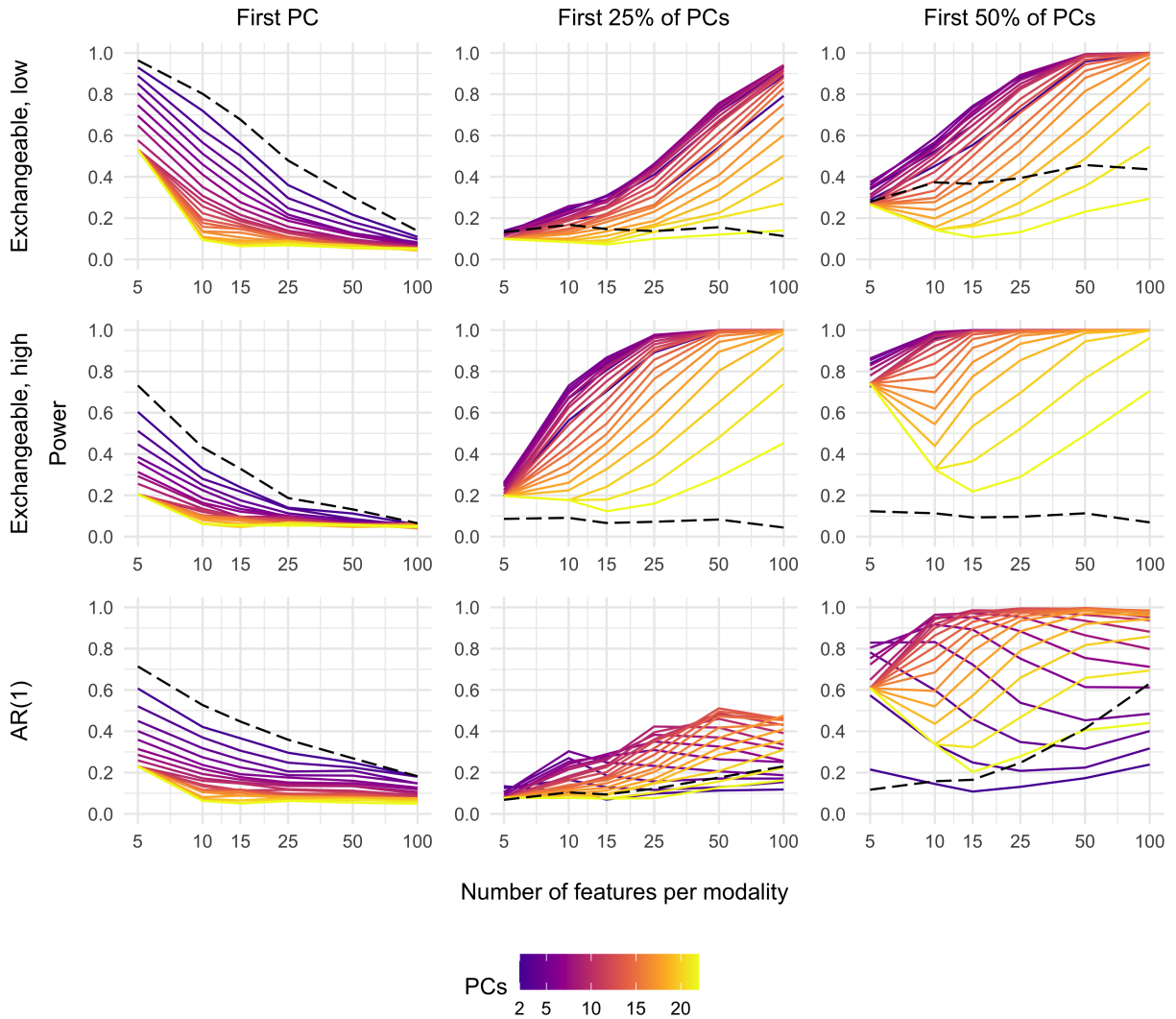
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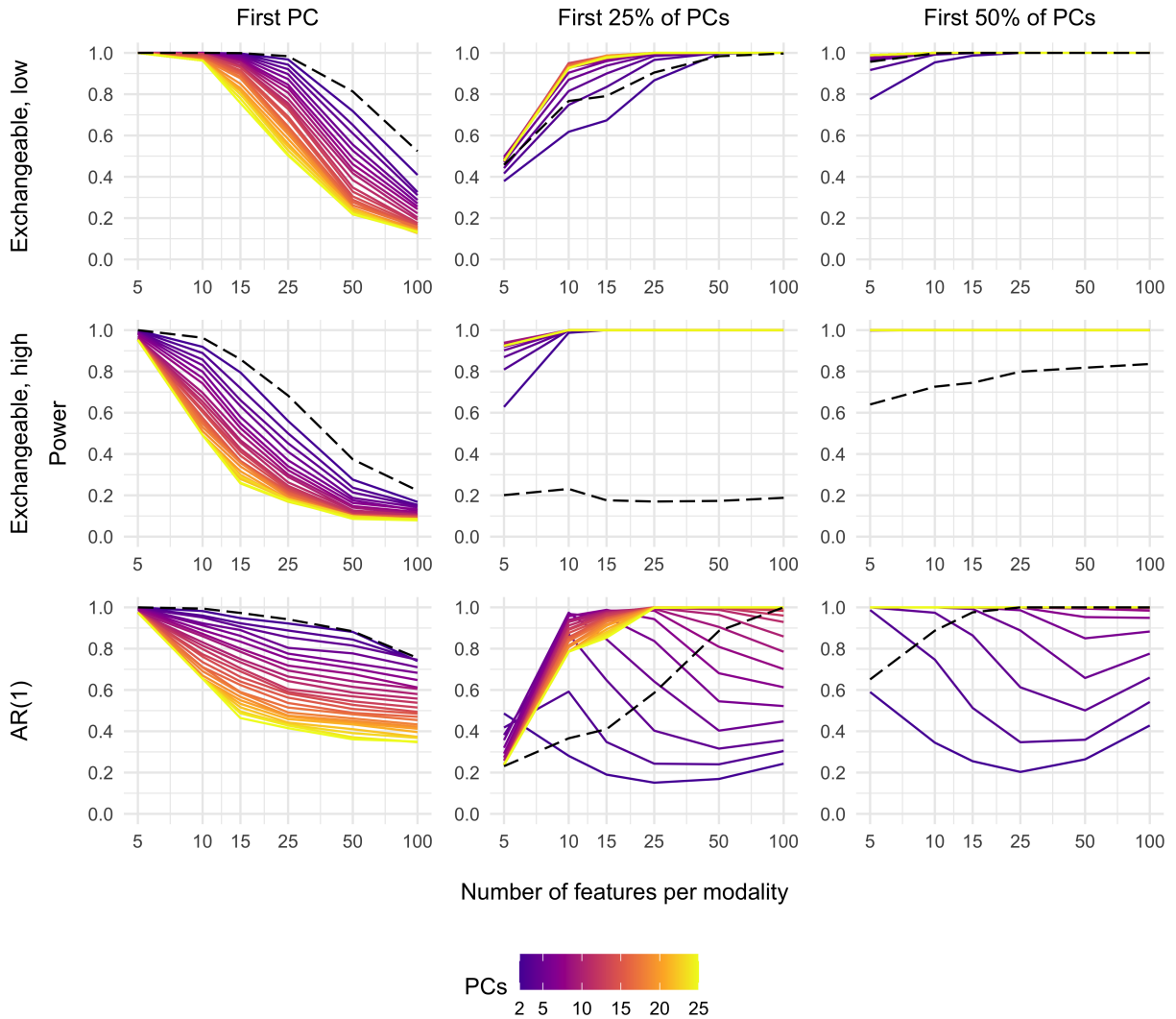
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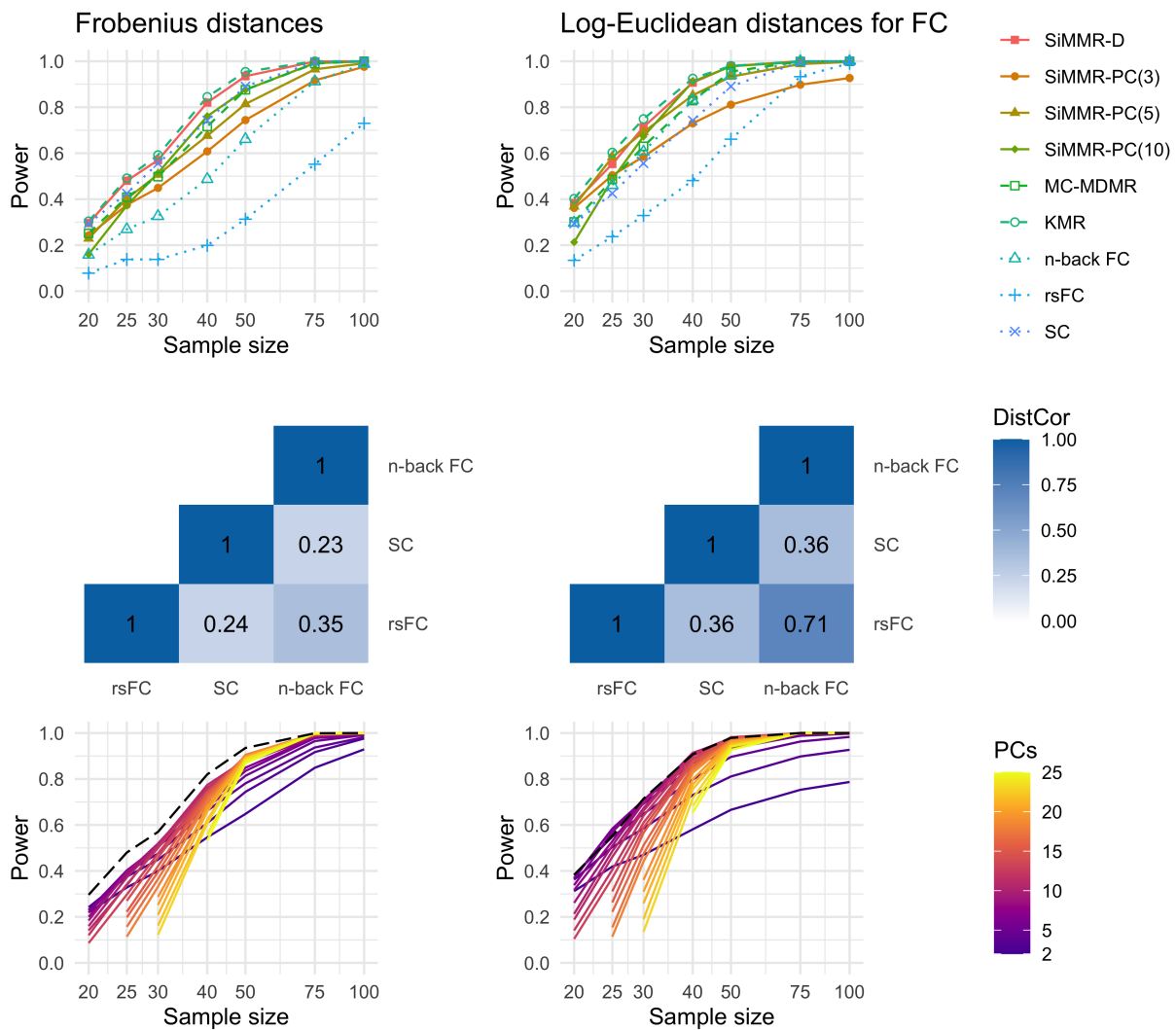




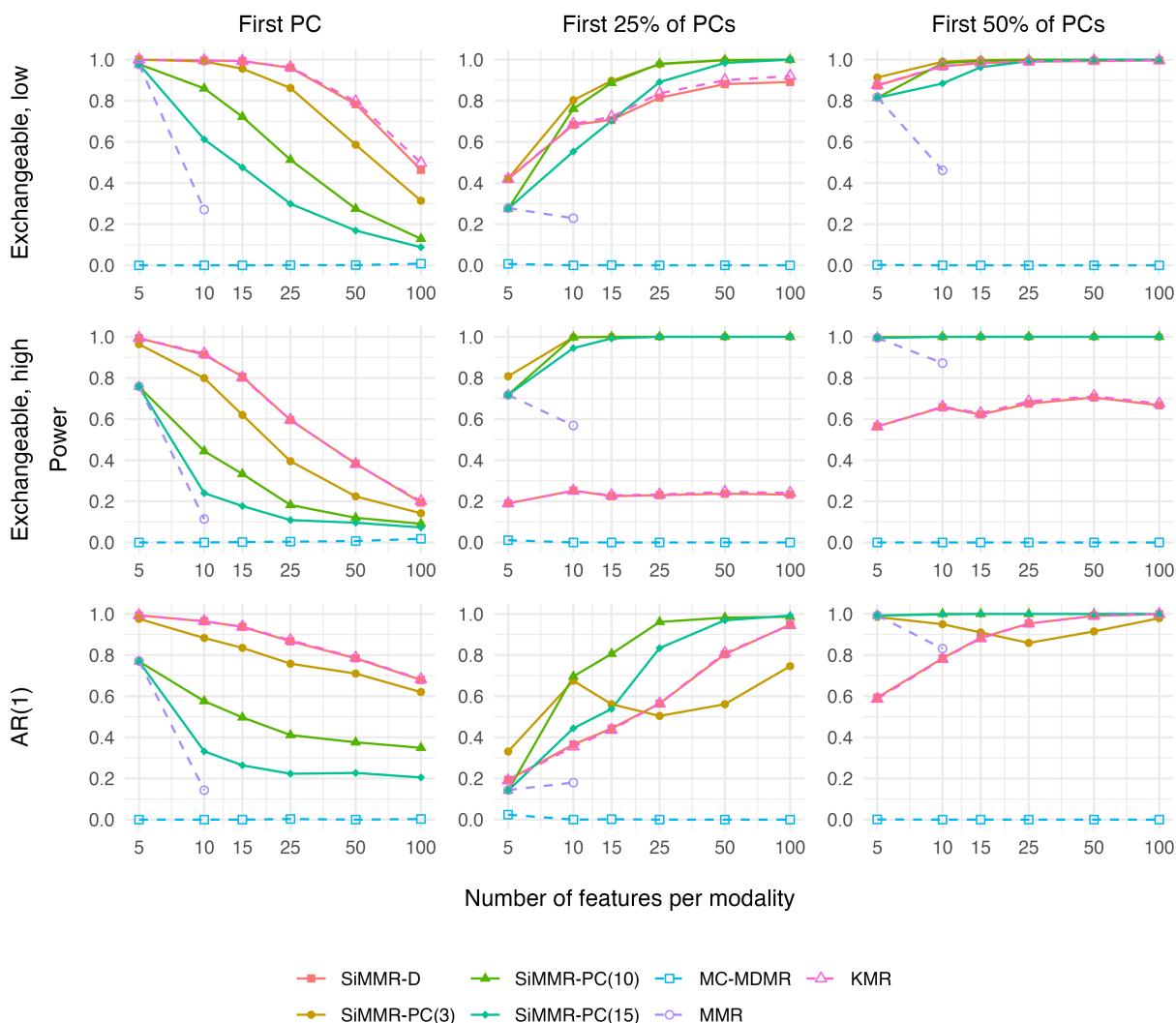
Supplementary Figure 2: **SiMMR-PC** power results in simulations with exchangeable and AR1 correlation structures with a sample size of 25. Results are shown for SiMMR-PC with PCs ranging from 2 to 25. The dashed line represents the power of SiMMR-D for comparison.



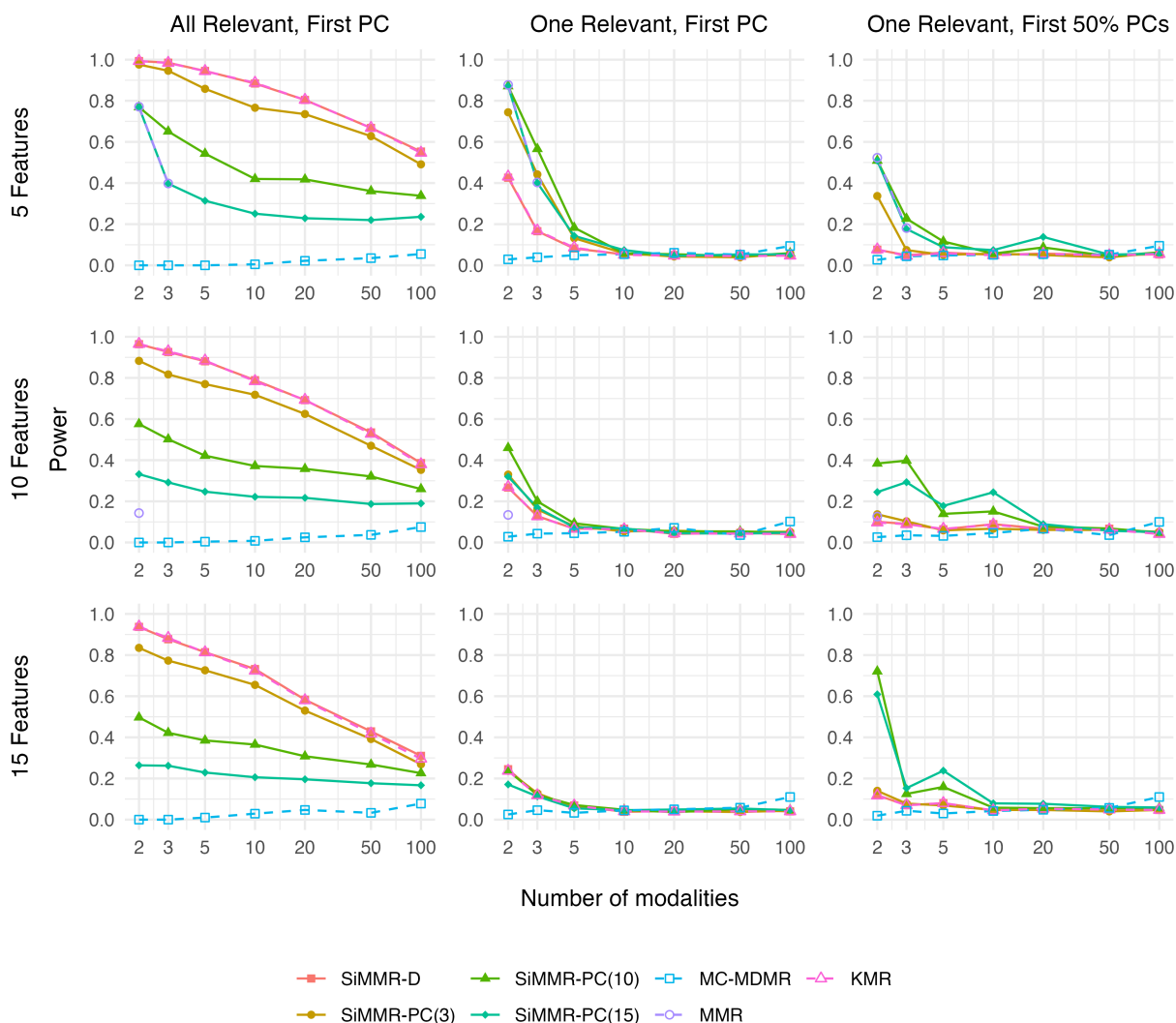
Supplementary Figure 3: **SiMMR-PC** power results in simulations with **exchangeable** and **AR1** correlation structures with a sample size of **100**. Results are shown for SiMMR-PC with PCs ranging from 2 to 25. The dashed line represents the power of SiMMR-D for comparison.



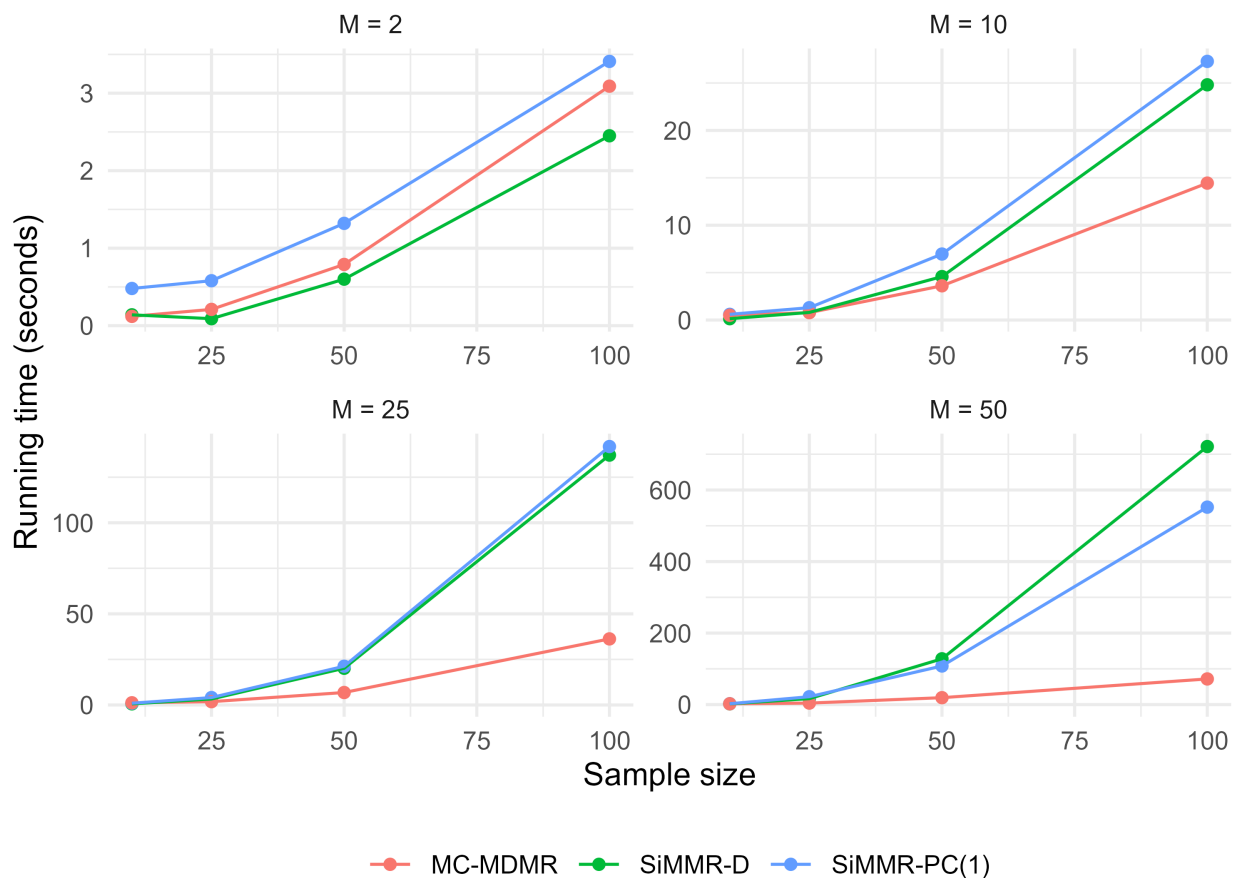
Supplementary Figure 4: **Comparison of Frobenius and log-Euclidean distances for PNC functional connectivity analyses.** Power for each test statistic and power across SiMMR-PC test statistics are shown for analyses using Frobenius versus log-Euclidean distances for rsFC and n-back FC dissimilarity matrices. Distance correlations are also displayed for the two choices of metric. Abbreviations: PNC, Philadelphia Neurodevelopmental Cohort; FC, functional connectivity; rsFC, resting-state functional connectivity; SC, structural connectivity.



Supplementary Figure 5: **Power results in continuous covariate simulations with exchangeable and AR(1) correlation structures for a sample size of 25.** Each trace represents a different test statistic. Different simulation settings are distinguished by correlation structure across rows and by rank of the binary covariate effect across columns. Exchangeable refers to an exchangeable correlation structure with low or high correlation and AR(1) refers to a first-order autoregressive structure. Abbreviations: MDMR, multivariate distance matrix regression; MC-MDMR, multiple MDMR statistics after Bonferroni correction; MMR, multivariate multiple regression using Pillai's trace.



Supplementary Figure 6: **Power results in AR(1) simulations where only one modality is relevant for a sample size of 25.** Each trace represents a different test statistic. Different simulation settings are distinguished by number of features per modality on the rows and simulation settings on the columns. All Relevant refers to the setting where every modality is related to the covariate of interest while One Relevant refers to a setting where only one modality is related, across all numbers of modalities. Abbreviations: MDMR, multivariate distance matrix regression; MC-MDMR, multiple MDMR statistics after Bonferroni correction; MMR, multivariate multiple regression using Pillai’s trace.



Supplementary Figure 7: **Running time in seconds for SiMMR test statistics and MC-MDMR across settings with exchangeable, high correlation.** Running time is reported for one simulated dataset in each setting. Settings vary in their number of observations (N) and number of modalities (M). All tests were run on a laptop with an Intel Core i9-13900H (24M Cache, up to 5.40 GHz) and 32 GB of RAM. Abbreviations: MDMR, multivariate distance matrix regression; MC-MDMR.

PC	N	M	Q	MMR	MCM	KMR	SiMMR-D	SiMMR-PC(3)	SiMMR-PC(10)	SiMMR-PC(15)
None	25	2	5	0.054	0.042	0.048	0.048	0.051	0.052	0.052
			25		0.048	0.043	0.044	0.051	0.049	0.045
			100		0.036	0.037	0.037	0.042	0.038	0.045
	10	2	5		0.032	0.05	0.048	0.033	0.035	0.037
			25		0.045	0.045	0.044	0.037	0.053	0.056
			100		0.05	0.042	0.041	0.04	0.041	0.054
	100	2	5	0.051	0.052	0.057	0.055	0.05	0.048	0.048
			25	0.053	0.045	0.046	0.042	0.047	0.048	0.046
		10	5	0.04	0.029	0.05	0.05	0.06	0.039	0.048
			25		0.051	0.046	0.042	0.042	0.044	0.042
		100	5		0.051	0.055	0.055	0.045	0.048	0.05
			100							
25%	25	2	5	0.078	0.081	0.065	0.068	0.125	0.077	0.077
			25		0.1	0.124	0.122	0.116	0.395	0.251
			100		0.151	0.23	0.23	0.154	0.391	0.454
		10	5		0.09	0.114	0.115	0.106	0.373	0.245
			25		0.134	0.295	0.305	0.202	0.395	0.493
			100		0.227	0.787	0.796	0.517	0.718	0.693
	100	2	5	0.241	0.291	0.231	0.231	0.418	0.237	0.237
			25	0.908	0.4	0.603	0.586	0.243	0.998	1
		10	5	0.9	0.409	0.576	0.568	0.222	1	1
			25		0.68	1	0.999	0.318	0.843	0.99
		100	5		1	1	1	0.92	0.999	1
			100							
50%	25	2	5	0.616	0.125	0.117	0.117	0.574	0.609	0.609
			25		0.167	0.238	0.247	0.208	0.983	0.971
			100		0.393	0.635	0.632	0.317	0.937	0.985
		10	5		0.136	0.238	0.233	0.187	0.982	0.975
			25		0.209	0.789	0.793	0.45	0.906	0.985
			100		0.554	1	1	0.954	0.999	0.996
	100	2	5	1	0.583	0.651	0.651	0.987	1	1
			25	1	0.989	1	1	0.347	1	1
		10	5	1	0.959	1	1	0.322	1	1
			25		0.995	1	1	0.648	1	1
		100	5		1	1	1	1	1	1
			100							

Supplementary Table 1: **Simulation results for the AR(1) correlation setting.** Rejection rates are shown across varying number of subjects (N), number of modalities (M), number of features per modality (Q), and number of principal components included in the binary covariate effect (PC). The highest power in each row is bolded.

PC	N	M	Q	Exchangeable, low correlation					Exchangeable, high correlation					AR(1) correlation					
				MMR	MCM	KMR	SiMMR-D	SiMMR-PC(3)	MMR	MCM	KMR	SiMMR-D	SiMMR-PC(3)	MMR	MCM	KMR	SiMMR-D	SiMMR-PC(3)	
None	25	2	5	0.038	0.044	0.05	0.049	0.05	0.038	0.051	0.047	0.046	0.049	0.038	0.051	0.045	0.045	0.043	
			100		0.032	0.047	0.038	0.05		0.039	0.041	0.043	0.049		0.038	0.041	0.043	0.051	
	10	5	5		0.06	0.053	0.047	0.047		0.053	0.048	0.045	0.043		0.052	0.039	0.039	0.042	
			100		0.05	0.064	0.048	0.042		0.038	0.054	0.048	0.04		0.052	0.052	0.057	0.046	
	100	2	5	0.053	0.054	0.05	0.05	0.057	0.053	0.052	0.051	0.051	0.056	0.053	0.054	0.053	0.052	0.052	0.056
			100		0.046	0.058	0.049	0.063		0.041	0.048	0.05	0.061		0.038	0.063	0.063	0.061	0.054
		10	5	5	0.057	0.048	0.039	0.039	0.052	0.057	0.047	0.044	0.043	0.051	0.057	0.054	0.042	0.041	0.047
				100		0.045	0.064	0.059	0.046		0.043	0.059	0.06	0.05		0.041	0.065	0.066	0.053
1	25	2	5	0.977	0	1	1	1	0.759	0	0.994	0.993	0.964	0.771	0	0.993	0.993	0.976	
			100		0.008	0.498	0.464	0.314		0.018	0.201	0.194	0.142		0.003	0.682	0.679	0.62	
	10	5	5		0.001	0.96	0.951	0.847		0.004	0.602	0.602	0.38		0.005	0.887	0.884	0.766	
			100		0.035	0.156	0.137	0.1		0.036	0.073	0.071	0.06		0.037	0.37	0.379	0.324	
	100	2	5	1	0	1	1	1	1	0	1	1	1	1	0	1	1	1	
			100		0	0.985	0.984	0.942		0.003	0.676	0.668	0.5		0	1	1	1	
		10	5	5	0.919	0	1	1	1	0.375	0	0.995	0.994	0.986	0.843	0	1	1	1
				100		0.006	0.506	0.48	0.318		0.023	0.204	0.2	0.129		0.009	0.991	0.99	0.943
25%	25	2	5	0.278	0.007	0.42	0.419	0.421	0.717	0.011	0.19	0.19	0.808	0.144	0.024	0.19	0.193	0.331	
			100		0	0.92	0.891	1		0	0.241	0.233	1		0	0.945	0.948	0.746	
	10	5	5		0.01	0.811	0.786	0.969		0.007	0.218	0.208	1		0.007	0.538	0.534	0.456	
			100		0	0.934	0.907	1		0	0.266	0.258	1		0	1	1	0.996	
	100	2	5	0.992	0	0.994	0.993	0.964	1	0.001	0.833	0.826	1	0.839	0	0.785	0.779	0.967	
			100		0	1	1	1		0	0.999	0.999	1		0	1	1	0.989	
		10	5	5	1	0.001	1	1	1	1	0	1	1	1	1	0	1	1	0.837
				100		0	1	1	1		0	1	1	1		0	1	1	1
50%	25	2	5	0.817	0.002	0.876	0.874	0.913	0.995	0	0.564	0.564	0.999	0.993	0.001	0.587	0.593	0.985	
			100		0	0.996	0.996	1		0	0.675	0.667	1		0	0.999	0.999	0.979	
	10	5	5		0.002	0.993	0.991	1		0.002	0.657	0.651	1		0.001	0.956	0.952	0.848	
			100		0	0.998	0.996	1		0	0.692	0.684	1		0	1	1	1	
	100	2	5	1	0	1	1	1	1	0	1	1	1	1	0	1	1	1	
			100		0	1	1	1		0	1	1	1		0	1	1	1	
		10	5	5	1	0	1	1	1	1	0	1	1	1	1	0	1	1	0.984
				100		0	1	1	1		0	1	1	1		0	1	1	1

Supplementary Table 2: **Continuous covariate simulation results across varying sample size, number of features, covariate effect, and correlation structure.** Rejection rates are shown across varying number of subjects (N), number of modalities (M), number of features per modality (Q), and number of principal components included in the binary covariate effect (PC). The highest power among tests within each simulation setting is bolded. Abbreviations: MMR, multivariate multiple regression using Pillai’s trace; MCM, multiple MDMR statistics after Bonferroni correction; KMR: kernel machine regression.

PC	N	M	Q	MMR	MCM	KMR	SiMMR-D	SiMMR-PC(3)	SiMMR-PC(10)	SiMMR-PC(15)
None	25	2	5	0.038	0.051	0.045	0.045	0.043	0.037	0.037
			25		0.044	0.044	0.045	0.056	0.046	0.046
			100		0.038	0.041	0.043	0.051	0.043	0.047
		10	5		0.052	0.039	0.039	0.042	0.045	0.043
			25		0.05	0.05	0.051	0.048	0.05	0.046
			100		0.052	0.052	0.057	0.046	0.057	0.049
	100	2	5	0.053	0.054	0.053	0.052	0.056	0.052	0.052
			25	0.058	0.043	0.06	0.058	0.064	0.056	0.061
			100		0.038	0.063	0.061	0.054	0.049	0.05
		10	5	0.057	0.054	0.042	0.041	0.047	0.05	0.057
			25		0.058	0.044	0.038	0.032	0.041	0.061
			100		0.041	0.065	0.066	0.053	0.061	0.065
25%	25	2	5	0.144	0.024	0.19	0.193	0.331	0.141	0.141
			25		0	0.565	0.564	0.504	0.961	0.834
			100		0	0.945	0.948	0.746	0.986	0.992
		10	5		0.007	0.538	0.534	0.456	0.951	0.816
			25		0	0.972	0.971	0.858	0.977	0.989
			100		0	1	1	0.996	1	1
	100	2	5	0.839	0	0.785	0.779	0.967	0.836	0.836
			25	1	0	1	1	0.9	1	1
			100		0	1	1	0.989	1	1
		10	5	1	0	1	1	0.837	1	1
			25		0	1	1	0.992	1	1
			100		0	1	1	1	1	1
50%	25	2	5	0.993	0.001	0.587	0.593	0.985	0.992	0.992
			25		0	0.953	0.953	0.859	1	1
			100		0	0.999	0.999	0.979	1	1
		10	5		0.001	0.956	0.952	0.848	1	1
			25		0	1	1	0.993	1	1
			100		0	1	1	1	1	1
	100	2	5	1	0	1	1	1	1	1
			25	1	0	1	1	0.992	1	1
			100		0	1	1	1	1	1
		10	5	1	0	1	1	0.984	1	1
			25		0	1	1	1	1	1
			100		0	1	1	1	1	1

Supplementary Table 3: **Continuous covariate simulation results for the AR(1) correlation setting.** Rejection rates are shown across varying number of subjects (N), number of modalities (M), number of features per modality (Q), and number of principal components included in the binary covariate effect (PC). The highest power in each row is bolded.