

**SupplementaryMaterial S5.**

Average cell subtype-wise performance results for cell subtype predictions of moSCminer with its variant and the baseline methods based on 5-fold cross-validation.

GSE136718	Cell subtype	moSCminer (omics-attn)	moSCminer (no-attn)	RF	SVM	LR	NB
Accuracy	2cell	<b>1.000</b>	<b>1.000</b>	<b>1.000</b>	<b>1.000</b>	<b>1.000</b>	0.971
	4cell	<b>0.991</b>	0.962	0.971	0.976	0.971	0.924
	8cell	0.995	0.995	<b>1.000</b>	0.995	0.991	0.910
	ICM	<b>1.000</b>	<b>1.000</b>	<b>1.000</b>	<b>1.000</b>	0.995	0.929
	Late4cell	<b>0.991</b>	0.962	0.971	0.971	0.967	0.933
	Morula	0.995	0.995	<b>1.000</b>	<b>1.000</b>	0.995	0.943
	TE	<b>1.000</b>	<b>1.000</b>	<b>1.000</b>	<b>1.000</b>	0.995	0.924
	Zygote	<b>1.000</b>	<b>1.000</b>	<b>1.000</b>	<b>1.000</b>	<b>1.000</b>	0.981
F1-score	2cell	<b>1.000</b>	<b>1.000</b>	<b>1.000</b>	<b>1.000</b>	<b>1.000</b>	0.870
	4cell	<b>0.977</b>	0.906	0.930	0.937	0.928	0.808
	8cell	0.960	0.987	<b>1.000</b>	0.985	0.973	0.736
	ICM	<b>1.000</b>	<b>1.000</b>	<b>1.000</b>	<b>1.000</b>	0.987	0.784
	Late4cell	<b>0.953</b>	0.780	0.847	0.825	0.798	0.565
	Morula	0.982	0.978	<b>1.000</b>	<b>1.000</b>	0.978	0.671
	TE	<b>1.000</b>	<b>1.000</b>	<b>1.000</b>	<b>1.000</b>	0.978	0.576
	Zygote	<b>1.000</b>	<b>1.000</b>	<b>1.000</b>	<b>1.000</b>	<b>1.000</b>	0.820
ROC-AUC	2cell	<b>1.000</b>	<b>1.000</b>	<b>1.000</b>	<b>1.000</b>	<b>1.000</b>	0.967
	4cell	<b>0.986</b>	0.952	0.982	0.985	0.982	0.921
	8cell	0.967	0.988	<b>1.000</b>	0.997	0.994	0.900
	ICM	<b>1.000</b>	<b>1.000</b>	<b>1.000</b>	<b>1.000</b>	0.997	0.869
	Late4cell	<b>0.981</b>	0.894	0.879	0.858	0.842	0.760
	Morula	0.997	0.997	<b>1.000</b>	<b>1.000</b>	0.980	0.789
	TE	<b>1.000</b>	<b>1.000</b>	<b>1.000</b>	<b>1.000</b>	0.980	0.819
	Zygote	<b>1.000</b>	<b>1.000</b>	<b>1.000</b>	<b>1.000</b>	<b>1.000</b>	0.867

GSE154762	Cell subtype	moSCminer (omics-attn)	moSCminer (no-attn)	RF	SVM	LR	NB
Accuracy	FGO	<b>0.993</b>	0.984	0.992	0.989	0.984	0.922
	GO1	<b>0.994</b>	0.971	0.986	0.983	0.972	0.940
	GO2	<b>0.992</b>	0.971	0.983	0.982	0.971	0.950
	Granulosa	0.991	0.984	0.990	<b>0.994</b>	0.993	0.912
	Immune	0.997	0.999	0.999	<b>1.000</b>	<b>1.000</b>	0.980
	MI	<b>0.982</b>	0.945	0.976	0.961	0.949	0.859
	MII	<b>0.989</b>	0.954	0.981	0.971	0.963	0.891
	StromaC1	<b>0.998</b>	0.912	0.953	0.920	0.911	0.797
StromaC2	<b>0.993</b>	0.911	0.954	0.923	0.909	0.786	
F1-score	FGO	<b>0.953</b>	0.911	0.965	0.932	0.907	0.560
	GO1	<b>0.922</b>	0.712	0.861	0.829	0.741	0.537
	GO2	<b>0.870</b>	0.600	0.741	0.740	0.607	0.504
	Granulosa	0.954	0.910	0.941	<b>0.973</b>	0.963	0.613
	Immune	0.978	0.978	0.921	<b>1.000</b>	<b>1.000</b>	0.160
	MI	<b>0.950</b>	0.844	0.928	0.884	0.854	0.544
	MII	<b>0.942</b>	0.768	0.898	0.848	0.808	0.505
	StromaC1	<b>0.995</b>	0.790	0.890	0.808	0.784	0.468
StromaC2	<b>0.982</b>	0.780	0.882	0.810	0.781	0.457	
ROC-AUC	FGO	<b>0.986</b>	0.944	0.961	0.949	0.944	0.763
	GO1	<b>0.981</b>	0.913	0.965	0.946	0.919	0.874
	GO2	<b>0.946</b>	0.754	0.811	0.840	0.760	0.765
	Granulosa	0.956	0.926	0.980	<b>0.991</b>	0.968	0.818
	Immune	0.999	0.999	0.930	<b>1.000</b>	<b>1.000</b>	0.550
	MI	<b>0.984</b>	0.914	0.963	0.936	0.920	0.715
	MII	<b>0.951</b>	0.873	0.933	0.914	0.885	0.744
	StromaC1	<b>0.999</b>	0.872	0.936	0.882	0.865	0.661
StromaC2	<b>0.992</b>	0.859	0.918	0.877	0.863	0.661	

GSE140203	Cell subtype	moSCminer (omics-attn)	moSCminer (no-attn)	RF	SVM	LR	NB
Accuracy	Basal	<b>0.998</b>	0.966	0.994	0.927	0.953	0.802
	Dermal Fibroblast	<b>0.999</b>	0.994	0.997	0.993	0.995	0.971
	Dermal Papilla	<b>0.998</b>	0.995	0.996	0.995	0.995	0.969
	Dermal Sheath	<b>0.999</b>	0.997	0.996	0.996	0.996	0.966
	Endothelial	<b>0.999</b>	0.994	0.998	0.993	0.993	0.976
	Granular	<b>0.999</b>	0.993	0.995	0.992	0.992	0.970
	Hair Shaft-cuticle-cortex	<b>0.998</b>	0.987	0.997	0.983	0.986	0.972
	IRS	<b>0.999</b>	0.989	0.997	0.985	0.987	0.969
	Infundibulum	<b>0.997</b>	0.964	0.984	0.947	0.958	0.887
	Isthmus	<b>0.999</b>	0.991	0.998	0.986	0.989	0.919
	K6+ Bulge Companion Layer	<b>0.999</b>	0.994	0.997	0.991	0.993	0.982
	Macrophage DC	<b>1.000</b>	0.997	0.998	0.995	0.996	0.982
	Medulla	<b>0.998</b>	0.985	0.996	0.982	0.984	0.964
	Melanocyte	<b>1.000</b>	0.999	0.998	0.999	0.999	0.989
	ORS	<b>0.999</b>	0.993	0.998	0.992	0.992	0.979
	Schwann Cell	<b>0.999</b>	0.998	0.999	0.996	0.997	0.852
	Sebaceous Gland	<b>1.000</b>	0.999	0.999	0.998	0.998	0.978
	Spinous	<b>0.997</b>	0.971	0.994	0.955	0.966	0.910
	TAC-1	<b>0.995</b>	0.959	0.972	0.944	0.952	0.908
	TAC-2	<b>0.998</b>	0.981	0.995	0.979	0.980	0.969
ahighCD34+ bulge	<b>0.999</b>	0.986	0.998	0.978	0.983	0.906	
alowCD34+ bulge	<b>0.998</b>	0.981	0.997	0.975	0.978	0.943	
F1-score	Basal	<b>0.995</b>	0.929	0.987	0.858	0.905	0.651
	Dermal Fibroblast	<b>0.980</b>	0.918	0.955	0.898	0.922	0.286
	Dermal Papilla	<b>0.968</b>	0.884	0.909	0.884	0.892	0.281
	Dermal Sheath	<b>0.945</b>	0.873	0.804	0.834	0.845	0.126
	Endothelial	<b>0.976</b>	0.901	0.958	0.869	0.877	0.330
	Granular	<b>0.924</b>	0.503	0.576	0.373	0.391	0.105
	Hair Shaft-cuticle-cortex	<b>0.978</b>	0.822	0.953	0.769	0.806	0.578
	IRS	<b>0.969</b>	0.726	0.909	0.635	0.676	0.397
	Infundibulum	<b>0.987</b>	0.865	0.941	0.801	0.841	0.330
	Isthmus	<b>0.979</b>	0.775	0.941	0.629	0.716	0.228
	K6+ Bulge Companion Layer	<b>0.969</b>	0.811	0.903	0.689	0.763	0.273
	Macrophage DC	<b>0.977</b>	0.805	0.828	0.625	0.728	0.075
	Medulla	<b>0.967</b>	0.755	0.930	0.671	0.718	0.162
	Melanocyte	<b>0.961</b>	0.900	0.837	0.884	0.873	0.378
	ORS	<b>0.982</b>	0.896	0.959	0.877	0.878	0.539
	Schwann Cell	<b>0.941</b>	0.724	0.823	0.456	0.542	0.035
	Sebaceous Gland	<b>0.992</b>	0.887	0.847	0.839	0.851	0.273
	Spinous	<b>0.987</b>	0.853	0.972	0.763	0.823	0.195
	TAC-1	<b>0.978</b>	0.804	0.881	0.742	0.777	0.285
	TAC-2	<b>0.965</b>	0.685	0.919	0.615	0.648	0.283
ahighCD34+ bulge	<b>0.991</b>	0.853	0.973	0.767	0.814	0.449	
alowCD34+ bulge	<b>0.985</b>	0.836	0.973	0.788	0.811	0.107	
ROC-AUC	Basal	<b>0.996</b>	0.955	0.989	0.919	0.946	0.789
	Dermal Fibroblast	<b>0.988</b>	0.971	0.985	0.950	0.959	0.585
	Dermal Papilla	<b>0.983</b>	0.928	0.968	0.930	0.930	0.616
	Dermal Sheath	<b>0.965</b>	0.919	0.837	0.890	0.896	0.570
	Endothelial	<b>0.990</b>	0.946	0.974	0.917	0.933	0.602
	Granular	<b>0.967</b>	0.698	0.703	0.629	0.648	0.585
	Hair Shaft-cuticle-cortex	<b>0.988</b>	0.909	0.955	0.879	0.897	0.760
	IRS	<b>0.982</b>	0.860	0.917	0.809	0.818	0.736
	Infundibulum	<b>0.994</b>	0.934	0.987	0.900	0.920	0.602
	Isthmus	<b>0.984</b>	0.876	0.944	0.785	0.830	0.736
	K6+ Bulge Companion Layer	<b>0.975</b>	0.910	0.912	0.812	0.863	0.607
	Macrophage DC	<b>0.978</b>	0.865	0.855	0.747	0.830	0.570
	Medulla	<b>0.983</b>	0.867	0.950	0.804	0.842	0.552
	Melanocyte	<b>0.967</b>	0.927	0.861	0.902	0.907	0.784
	ORS	<b>0.988</b>	0.945	0.961	0.925	0.933	0.694
	Schwann Cell	<b>0.953</b>	0.806	0.853	0.658	0.712	0.665
	Sebaceous Gland	<b>0.992</b>	0.937	0.868	0.888	0.894	0.818

Spinous	<b>0.994</b>	0.918	0.990	0.861	0.902	0.554
TAC-1	<b>0.992</b>	0.893	0.982	0.871	0.886	0.586
TAC-2	<b>0.980</b>	0.824	0.925	0.770	0.795	0.597
ahighCD34+ bulge	<b>0.993</b>	0.927	0.974	0.868	0.893	0.844
alowCD34+ bulge	<b>0.994</b>	0.916	0.978	0.888	0.906	0.529

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