

## Supplemental Material

**Table S1.** List of pretrained deep networks used.

Modelname	Pretraining data	Number of parameters	Size of feature vector
<b>radimagenet.resnet50</b>	Medical data	23,59	2048
<b>radimagenet.densenet121</b>	Medical data	7,04	1024
<b>radimagenet.inceptionV3</b>	Medical data	21,8	2048
<b>radimagenet.IRV2</b>	Medical data	54,33	1536
<b>medicalImagenet.resnet10</b>	Medical data	14,36	512
<b>medicalImagenet.resnet18</b>	Medical data	32,99	512
<b>medicalImagenet.resnet34</b>	Medical data	63,3	512
<b>medicalImagenet.resnet50</b>	Medical data	46,16	2048
<b>convnext-v2-large_fcmae-in21k-pre_3rdparty_in1k</b>	ImageNet-1K	197,96	1536
<b>deit3-huge-p14_in21k-pre_3rdparty_in1k</b>	ImageNet-1K	632,13	1280
<b>efficientnet-b7_3rdparty-ra-noisystudent_in1k</b>	ImageNet-1K	66,35	80
<b>efficientnetv2-1_in21k-pre_3rdparty_in1k</b>	ImageNet-1K	118,52	96
<b>simclr_resnet50_16xb256-coslr-200e_in1k</b>	Self-supervised	27,97	2048
<b>simsiam_resnet50_8xb32-coslr-200e_in1k</b>	Self-supervised	38,2	2048
<b>mocov3_resnet50_8xb512-amp-coslr-300e_in1k</b>	Self-supervised	68,01	2048
<b>barlowtwins_resnet50_8xb256-coslr-300e_in1k</b>	Self-supervised	174,54	2048
<b>resnet34_8xb32_in1k</b>	ImageNet-1K	2,18	512
<b>vgg16bn_8xb32_in1k</b>	ImageNet-1K	138,37	1000
<b>densenet161_3rdparty_in1k</b>	ImageNet-1K	28,68	2208
<b>efficientnet-b2_3rdparty_8xb32_in1k</b>	ImageNet-1K	9,11	48

**Table S2.** AUC of the models using hand-crafted features.

Model	C4KC-KiTS	CRLM	Desmoid	GBM	GIST	HN	I-SPY1	Lipo	Liver	Melanoma	Mean AUC	ΔAUC
<b>Hand-crafted, 2-D, All</b>	0.761	<b>0.79</b>	0.899	0.722	0.817	0.902	0.68	0.883	0.801	0.681	0.794	0.005
<b>Hand-crafted, 2-D, No morph</b>	0.756	0.778	0.891	0.741	<b>0.819</b>	0.902	0.68	<b>0.885</b>	<b>0.803</b>	0.681	0.793	0.005
<b>Hand-crafted, 3-D, No morph</b>	<b>0.768</b>	0.749	0.898	0.772	0.764	<b>0.914</b>	0.671	0.876	0.782	<b>0.719</b>	0.791	0.002
<b>Hand-crafted, 3-D, All</b>	<b>0.768</b>	0.749	<b>0.908</b>	<b>0.783</b>	0.78	0.903	0.671	0.883	0.774	0.669	0.789	0
<b>Hand-crafted, 3-D, Morph</b>	0.58	0.742	0.807	0.694	0.726	0.849	<b>0.681</b>	0.884	0.676	0.589	0.723	-0.066

Models with “All” used all available hand-crafted features, while those with “No morph” used all but morphological features. The models with “Morph” used only morphological features. ΔAUC denotes the mean difference to the standard “Hand-crafted, 3-D, All” model. Models with the highest AUC for each dataset were marked in bold.

**Table S3.** AUC of the models using deep features.

Model	C4KC-KiTS	CRLM	Desmoid	GBM	GIST	HN	I-SPY1	Lipo	Liver	Melanoma	Mean AUC	ΔAUC
<b>ConvNeXt V2, large</b>	0.661	<b>0.821</b>	<b>0.881</b>	0.781	0.742	0.902	0.661	<b>0.886</b>	0.793	0.623	0.775	-0.014
<b>SimSiam, ResNet-50</b>	0.664	0.671	0.829	<b>0.915</b>	0.689	0.911	0.732	0.846	0.794	0.614	0.766	-0.022
<b>SimCLR, ResNet-50</b>	0.682	0.714	0.859	0.772	0.701	0.908	0.688	0.822	<b>0.819</b>	0.649	0.761	-0.027
<b>DenseNet-161</b>	0.675	0.676	0.867	0.709	<b>0.757</b>	0.89	0.688	0.843	0.778	0.666	0.755	-0.034
<b>ResNet-34</b>	<b>0.711</b>	0.743	0.845	0.645	0.738	0.874	0.652	0.885	0.816	0.607	0.752	-0.037
<b>VGG-16</b>	0.625	0.667	0.872	0.725	0.703	0.85	0.68	0.857	0.786	0.742	0.751	-0.038
<b>BarlowTwins, ResNet-50</b>	0.692	0.672	0.82	0.726	0.697	0.896	0.755	0.826	0.786	0.631	0.75	-0.039
<b>DeiT III huge</b>	0.646	0.689	0.761	0.785	0.738	0.864	<b>0.788</b>	0.871	0.702	0.644	0.749	-0.04
<b>EfficientNet-B2</b>	0.657	0.705	0.788	0.734	0.711	0.831	0.765	0.821	0.773	0.654	0.744	-0.045
<b>MoCo V3, ResNet-50</b>	0.67	0.734	0.872	0.651	0.685	<b>0.913</b>	0.674	0.827	0.779	0.6	0.74	-0.048
<b>EfficientNet-B7</b>	0.664	0.722	0.818	0.678	0.687	0.886	0.669	0.835	0.703	0.694	0.736	-0.053
<b>EfficientNet-V2, large</b>	0.671	0.668	0.816	0.725	0.706	0.856	0.707	0.762	0.749	0.656	0.732	-0.057
<b>MedicalNet, ResNet-10</b>	0.688	0.652	0.749	0.819	0.736	0.812	0.645	0.758	0.726	0.712	0.73	-0.059
<b>MedicalNet, ResNet-50</b>	0.712	0.59	0.721	0.803	0.697	0.816	0.647	0.693	0.737	<b>0.775</b>	0.719	-0.07
<b>RadImageNet, InceptionResNet-V2</b>	0.61	0.669	0.743	0.756	0.689	0.744	0.663	0.743	0.645	<b>0.775</b>	0.704	-0.085
<b>RadImageNet, DenseNet-121</b>	0.61	0.684	0.746	0.699	0.706	0.751	0.707	0.778	0.719	0.599	0.7	-0.089
<b>MedicalNet, ResNet-18</b>	0.679	0.565	0.756	0.726	0.664	0.823	0.65	0.735	0.719	0.677	0.7	-0.089
<b>RadImageNet, Inception-V3</b>	0.599	0.642	0.774	0.781	0.669	0.78	0.638	0.75	0.688	0.576	0.69	-0.099
<b>MedicalNet, ResNet-34</b>	0.7	0.569	0.736	0.724	0.67	0.778	0.645	0.743	0.667	0.648	0.688	-0.101
<b>RadImageNet, ResNet-50</b>	0.642	0.715	0.691	0.766	0.676	0.714	0.688	0.702	0.598	0.662	0.685	-0.103

ΔAUC denotes the mean difference to the standard “Hand-crafted, 3-D, All” model. Models with the highest AUC for each dataset were marked in bold.

**Table S4.** AUC of the models using deep features fused with morphological features.

Model	C4KC-KiTS	CRLM	Desmoid	GBM	GIST	HN	I-SPY1	Lipo	Liver	Melanoma	Mean AUC	ΔAUC
<b>ConvNeXt V2, large</b>	0.661	<b>0.792</b>	<b>0.882</b>	0.801	0.746	0.902	0.674	0.884	<b>0.818</b>	0.618	0.778	-0.011
<b>SimSiam, ResNet-50</b>	0.699	0.671	0.851	<b>0.915</b>	0.675	0.902	0.732	0.884	0.793	0.644	0.777	-0.012
<b>EfficientNet-B2</b>	0.652	0.794	0.847	0.839	0.717	0.864	0.71	0.884	0.758	0.665	0.773	-0.016
<b>SimCLR, ResNet-50</b>	0.682	0.703	0.872	0.768	0.722	0.91	0.698	0.884	0.807	0.611	0.766	-0.023
<b>DenseNet-161</b>	0.675	0.684	0.867	0.731	<b>0.758</b>	0.896	0.688	0.884	0.778	0.67	0.763	-0.026
<b>BarlowTwins, ResNet-50</b>	0.692	0.68	0.855	0.726	0.707	0.904	0.744	0.884	0.782	0.628	0.76	-0.029
<b>VGG-16</b>	0.628	0.688	0.877	0.784	0.712	0.847	0.64	0.884	0.795	0.723	0.758	-0.031
<b>EfficientNet-V2, large</b>	0.673	0.715	0.842	0.735	0.726	0.861	0.715	0.884	0.76	0.659	0.757	-0.032
<b>ResNet-34</b>	<b>0.715</b>	0.717	0.855	0.708	0.747	0.868	0.65	0.896	0.817	0.581	0.755	-0.033
<b>DeiT III huge</b>	0.646	0.71	0.773	0.798	0.741	0.865	<b>0.781</b>	<b>0.918</b>	0.678	0.644	0.755	-0.034
<b>MedicalNet, ResNet-50</b>	0.712	0.669	0.784	0.803	0.686	0.845	0.612	0.884	0.756	<b>0.776</b>	0.753	-0.036
<b>EfficientNet-B7</b>	0.664	0.742	0.842	0.714	0.725	0.875	0.663	0.884	0.713	0.695	0.752	-0.037
<b>MedicalNet, ResNet-10</b>	0.687	0.686	0.795	0.821	0.722	0.849	0.649	0.884	0.749	0.669	0.751	-0.038
<b>MoCo V3, ResNet-50</b>	0.67	0.72	0.876	0.651	0.699	<b>0.913</b>	0.681	0.884	0.779	0.618	0.749	-0.04
<b>RadImageNet, ResNet-50</b>	0.652	0.758	0.777	0.759	0.7	0.849	0.708	0.884	0.651	0.675	0.741	-0.048
<b>RadImageNet, InceptionResNet-V2</b>	0.666	0.66	0.781	0.752	0.706	0.841	0.673	0.888	0.654	0.775	0.74	-0.049
<b>MedicalNet, ResNet-18</b>	0.688	0.649	0.835	0.755	0.686	0.843	0.636	0.884	0.73	0.677	0.738	-0.051
<b>MedicalNet, ResNet-34</b>	0.7	0.633	0.818	0.738	0.697	0.849	0.644	0.884	0.727	0.648	0.734	-0.055
<b>RadImageNet, DenseNet-121</b>	0.602	0.685	0.77	0.689	0.748	0.836	0.664	0.887	0.709	0.606	0.72	-0.069
<b>RadImageNet, Inception-V3</b>	0.58	<b>0.674</b>	0.8	0.788	<b>0.682</b>	0.856	0.616	0.884	0.662	0.542	0.708	-0.08

ΔAUC denotes the mean difference to the standard “Hand-crafted, 3-D, All” model. Models with the highest AUC for each dataset were marked in bold.

**Table S5.** AUC of the models using deep features fused with hand-crafted features.

ModelName	C4KC-KiTS	CRLM	Desmoid	GBM	GIST	HN	I-SPY1	Lipo	Liver	Melanoma	Mean AUC	ΔAUC
<b>SimSiam, ResNet-50</b>	0.74	0.711	0.868	<b>0.915</b>	0.768	0.904	<b>0.704</b>	<b>0.933</b>	0.794	0.647	0.798	0.009
<b>EfficientNet-V2, large</b>	0.772	0.738	<b>0.925</b>	0.764	0.777	0.905	0.701	0.885	0.787	0.693	0.795	0.006
<b>EfficientNet-B2</b>	0.768	0.76	0.911	0.779	0.77	0.909	0.655	0.87	0.8	0.723	0.795	0.006
<b>MedicalNet, ResNet10</b>	0.754	0.749	0.891	0.819	0.78	0.896	0.649	0.873	0.779	0.73	0.792	0.003
<b>DeiT III huge</b>	0.759	0.753	0.852	0.754	<b>0.815</b>	0.882	0.701	0.884	0.788	0.718	0.791	0.002
<b>EfficientNet-B7</b>	0.771	0.772	0.923	0.754	0.762	0.906	0.676	0.873	0.768	0.699	0.79	0.001
<b>SimCLR, ResNet-50</b>	<b>0.794</b>	0.77	0.896	0.739	0.77	<b>0.911</b>	0.677	0.885	0.787	0.652	0.788	-0.001
<b>RadImageNet, ResNet50</b>	0.758	0.744	0.87	0.795	0.767	0.899	0.686	0.889	0.774	0.688	0.787	-0.002
<b>ConvNeXt V2, large</b>	0.743	<b>0.809</b>	0.911	0.751	0.781	0.902	0.656	0.882	0.789	0.643	0.787	-0.002
<b>MedicalNet, ResNet34</b>	0.771	0.767	0.883	0.781	0.782	0.899	0.671	0.885	0.766	0.657	0.786	-0.003
<b>ResNet-34</b>	0.763	0.74	0.895	0.765	0.791	0.889	0.643	0.886	<b>0.814</b>	0.644	0.783	-0.006
<b>BarlowTwins, ResNet-50</b>	0.737	0.749	0.889	0.749	0.785	0.886	0.701	0.875	0.785	0.674	0.783	-0.006
<b>VGG-16</b>	0.743	0.735	0.918	0.752	0.768	0.878	0.657	0.877	0.801	0.697	0.783	-0.006
<b>RadImageNet, InceptionResNet-V2</b>	0.758	0.742	0.876	0.701	0.761	0.899	0.683	0.873	0.774	<b>0.734</b>	0.78	-0.009
<b>MoCo V3, ResNet-50</b>	0.764	0.777	0.897	0.684	0.768	0.906	0.679	0.879	0.81	0.635	0.78	-0.009
<b>DenseNet-161</b>	0.78	0.735	0.909	0.697	0.795	0.89	0.669	0.884	0.803	0.631	0.779	-0.01
<b>RadImageNet, DenseNet-121</b>	0.745	0.784	0.866	0.749	0.784	0.899	0.664	0.874	0.764	0.662	0.779	-0.01
<b>MedicalNet, ResNet-18</b>	0.744	0.748	0.882	0.748	0.758	0.899	0.671	0.881	0.761	0.681	0.777	-0.012
<b>MedicalNet, ResNet-50</b>	0.739	0.725	0.866	0.747	0.775	0.897	0.619	0.876	0.795	0.712	0.775	-0.014
<b>RadImageNet, Inception-V3</b>	0.741	<b>0.749</b>	0.874	0.742	0.757	0.899	0.644	0.873	0.738	0.675	0.769	-0.02

ΔAUC denotes the mean difference to the standard “Hand-crafted, 3-D, All” model. Models with the highest AUC for each dataset were marked in bold.