

Electronic Supplementary Information

for

Bayesian graph convolutional network for reliable prediction of molecular properties with uncertainty quantification

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Korea

1. Training details

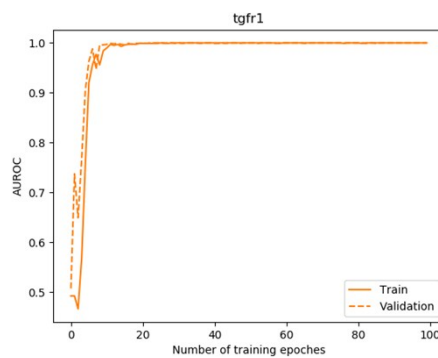
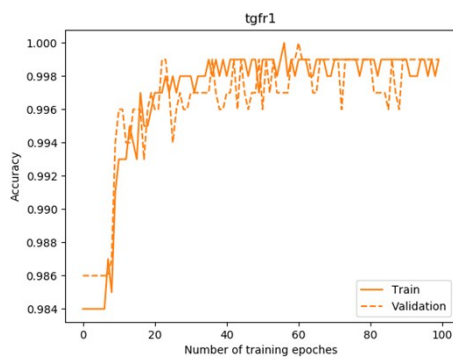
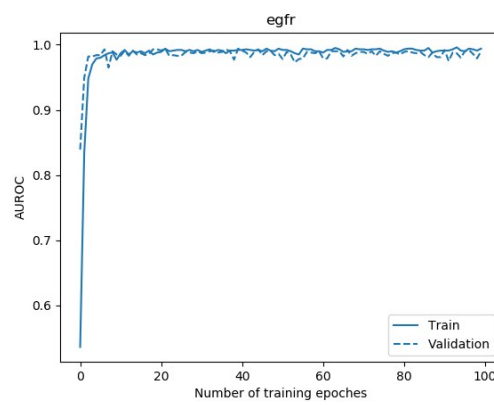
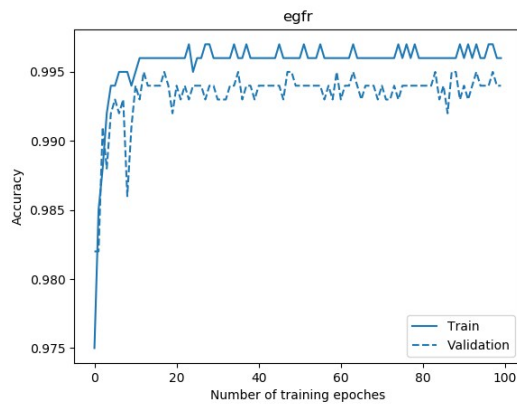
(1) The models in section 4.1

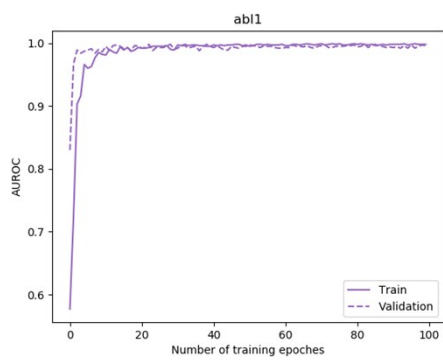
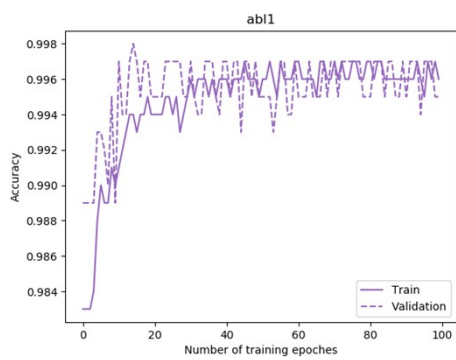
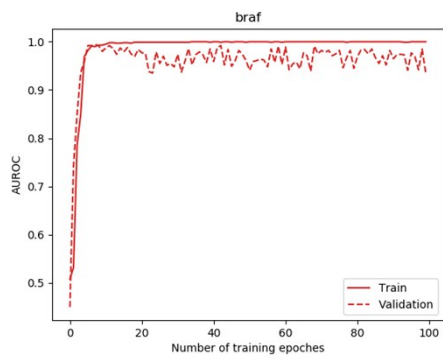
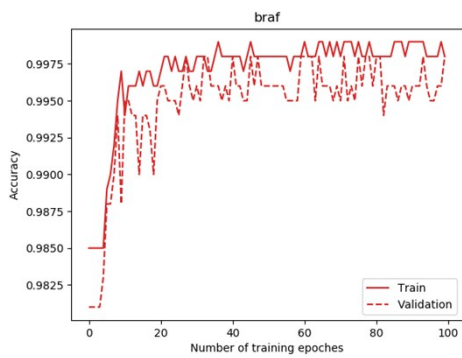
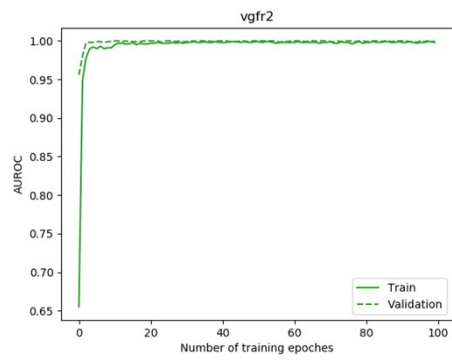
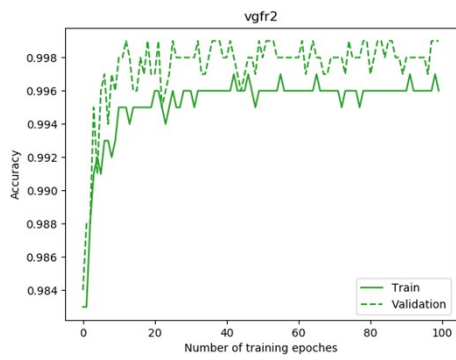
1) DUD-E dataset

* Number of training data and time for training

	Number of samples			Time for training (s)
	Train	Validation	Test	
egfr	25627	2847	7118	9887
tgfr1	6217	690	1726	2605
vgfr2	18259	2028	5071	7234
braf	7274	808	2020	3319
abl1	7872	874	2186	3581

* Accuracy & AUROC curve



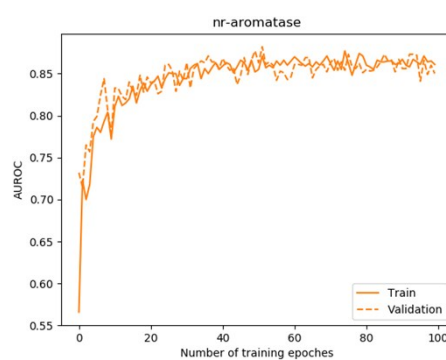
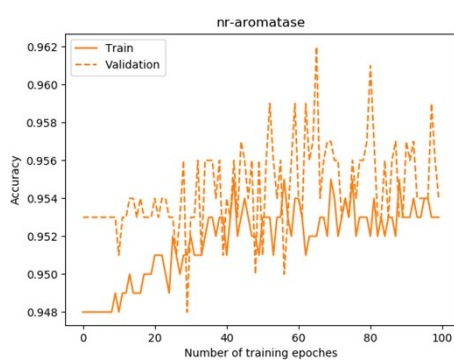
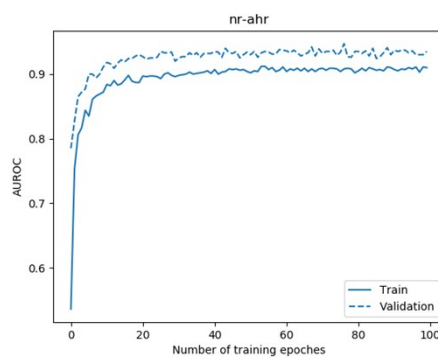
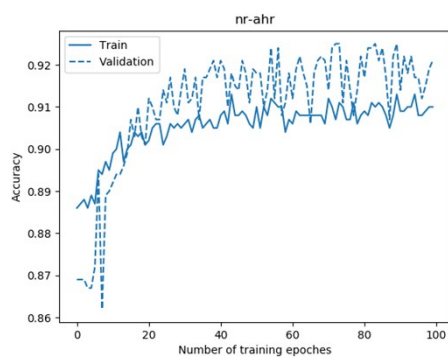


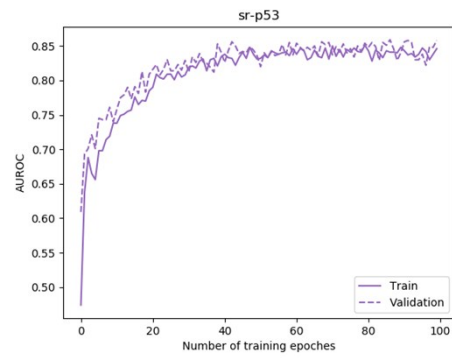
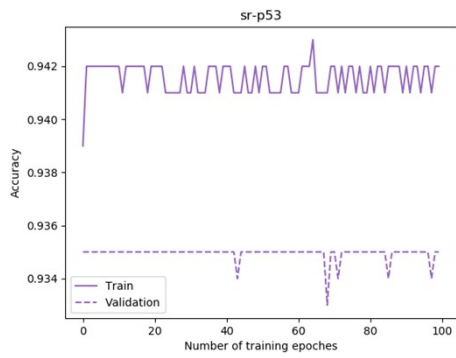
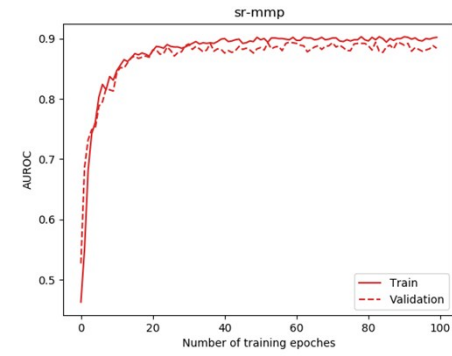
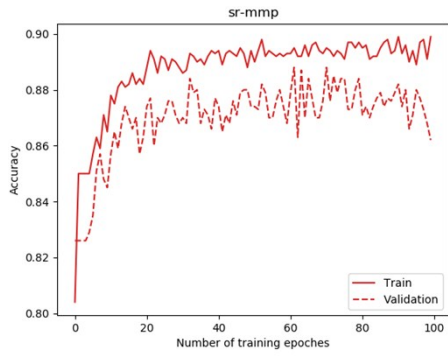
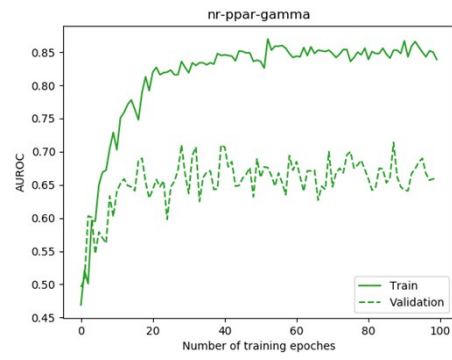
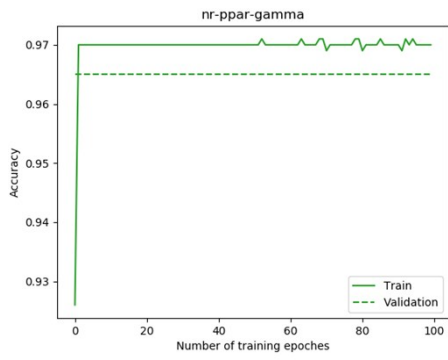
2) Tox-21 dataset

* Number of training data and time for training

	Number of samples			Time for training (s)
	Train	Validation	Test	
nr-ahr	6481	720	1800	2680
nr-aromatase	5706	634	1585	2358
nr-ppar-gamma	6489	721	1802	2644
sr-mmp	5801	644	1611	2439
sr-p53	6813	756	1892	2508

* Accuracy & AUROC curve



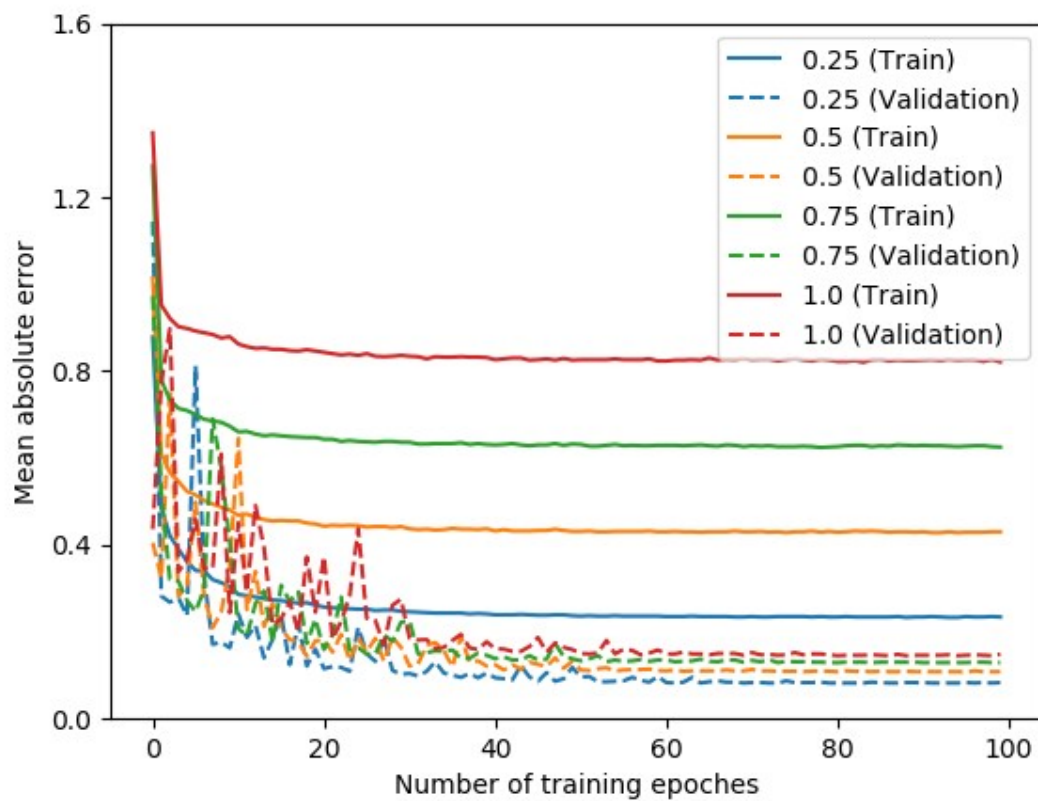


(2) The models in section 4.3

* Number of training data and time for training

	Number of samples			Time for training (s)
	Train	Validation	Test	
$\sigma^2 = 0.25, 0.5, 0.75, 1.0$	97287	10809	27023	$\sim 4 \times 10^4$

* Mean absolute error curve



Note that we adjusted noise only for the training samples. (Therefore, the mean absolute error of training curves are shifted.)

(3) The model in section 4.4

* Number of training data and time for training

	Number of samples			Time for training (s)
	Train	Validation	Test	
-	21585	2398	5995	8433

* Mean absolute error curve

