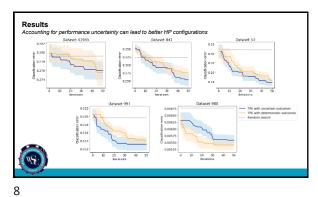


HPI			Cross validation protocol.						CD	Ž(y ▲		
max_iter	Z	U[10,1000]			Split 1	Fold 1	Fold 2	Fold 3	Fold 4	Fold 5	+y1	Г
neurons	Z	U[5, 1000]		New HP configuration	Split 2	Fold 1	Fold 2 Fold 2 Fold 2 Fold 2	Fold 3 Fold 3 Fold 3	Fold 4 Fold 4 Fold 4	Fold 5 Fold 5	• y <sub>21</sub> • y <sub>3</sub> • y <sub>4</sub> • y <sub>5</sub>	1 106 t
Ir init	init ℤ −U lo	g([10-6, 10-1])			Split 3	Fold 1						
h1		log([10 <sup>-7</sup> ,1])			Split 4	Fold 1						
h2		log([10 <sup>-7</sup> ,1])			Split 5					Fold 5		
	Classification			Training set				Validation	set			
Dataset	OpenMLID	Instances	Algorithms setting									
Balance scale	997	4	625	Initial design space						13	11d - 1 = 54	
Optdigits	980	64	5620	Replications to account for uncertainty (k in					5			
Stock	841	9	950	cross-validation protocol)								
Heart-statlog	53 13 270 Iterations							50				
Ilpd	41945	10	583	Candidates to sample per iteration $(n_c)$							2000	
				TPE(γ)							0.2	
				Macro	-replica	tions					10	



## Concluding remarks

- Basic idea: TPE algorithm adjustment to account for performance uncertainty
  Weighted KDE: probability that a given HP configuration is "good" or "bad"
- Our proposal outperforms the original TPE (final result and/or search speed)
  Interesting for settings with limited budget!
- Further fine-tuning required to get high-quality performance on datasets with probability of being good and bad very close (dataset 980)





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