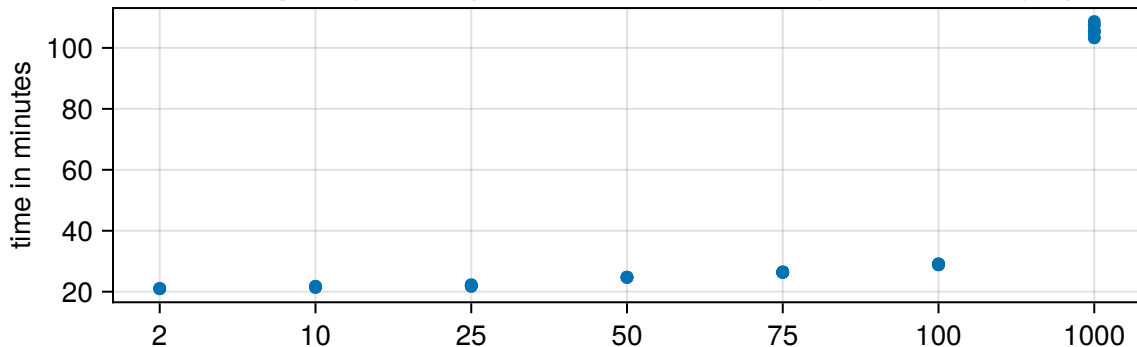


**Total computing times by number of nodes in the gene embedding layer**



**Average speed of training by number of nodes in the gene embedding layer**

