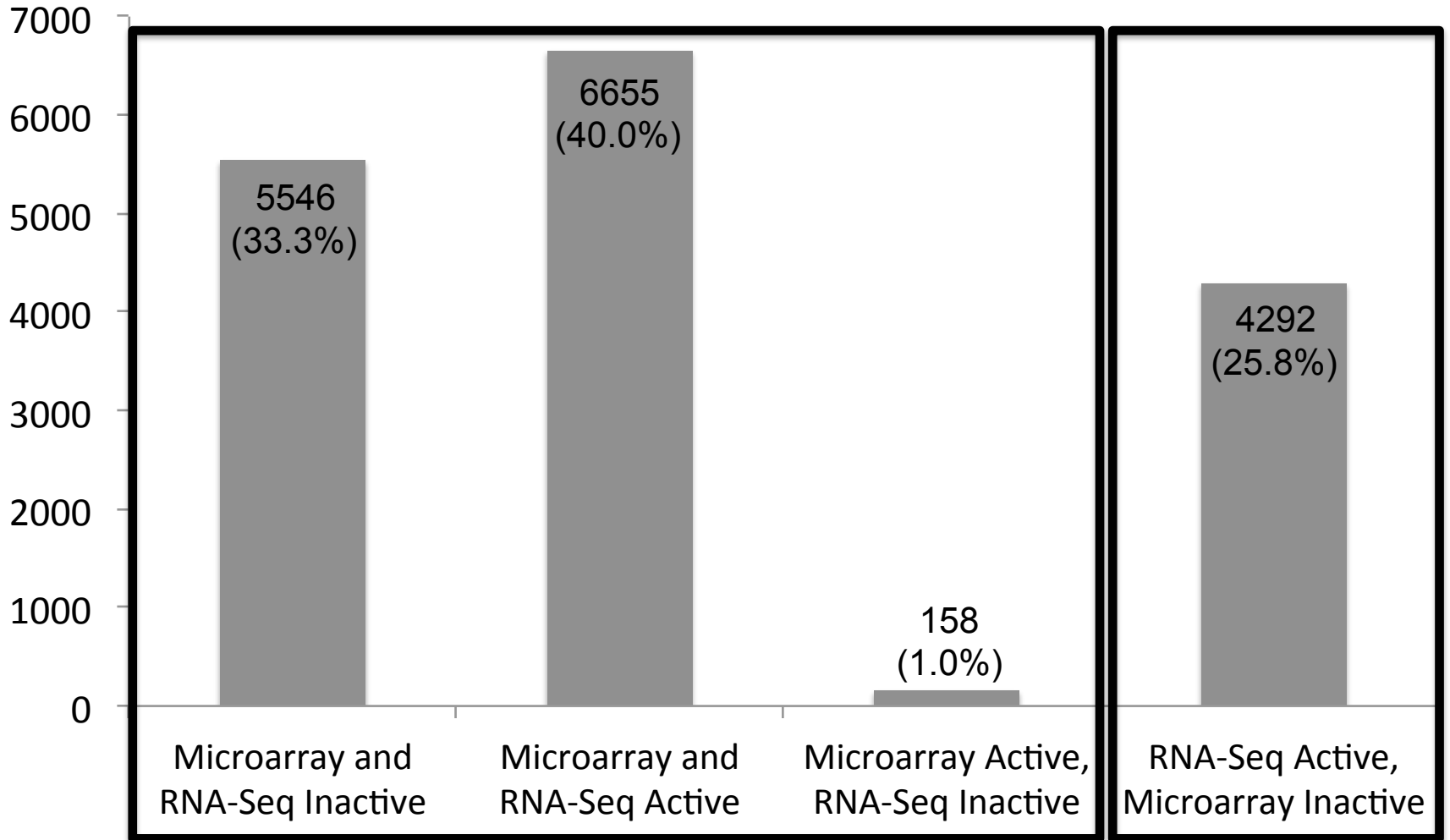


## Active and Inactive Genes Across Platforms



**Microarray active or  
RNA-Seq inactive subset:  
98.7% correspondence  
across platforms**

**Discordant due  
to increased  
sensitivity of  
RNA-Seq**