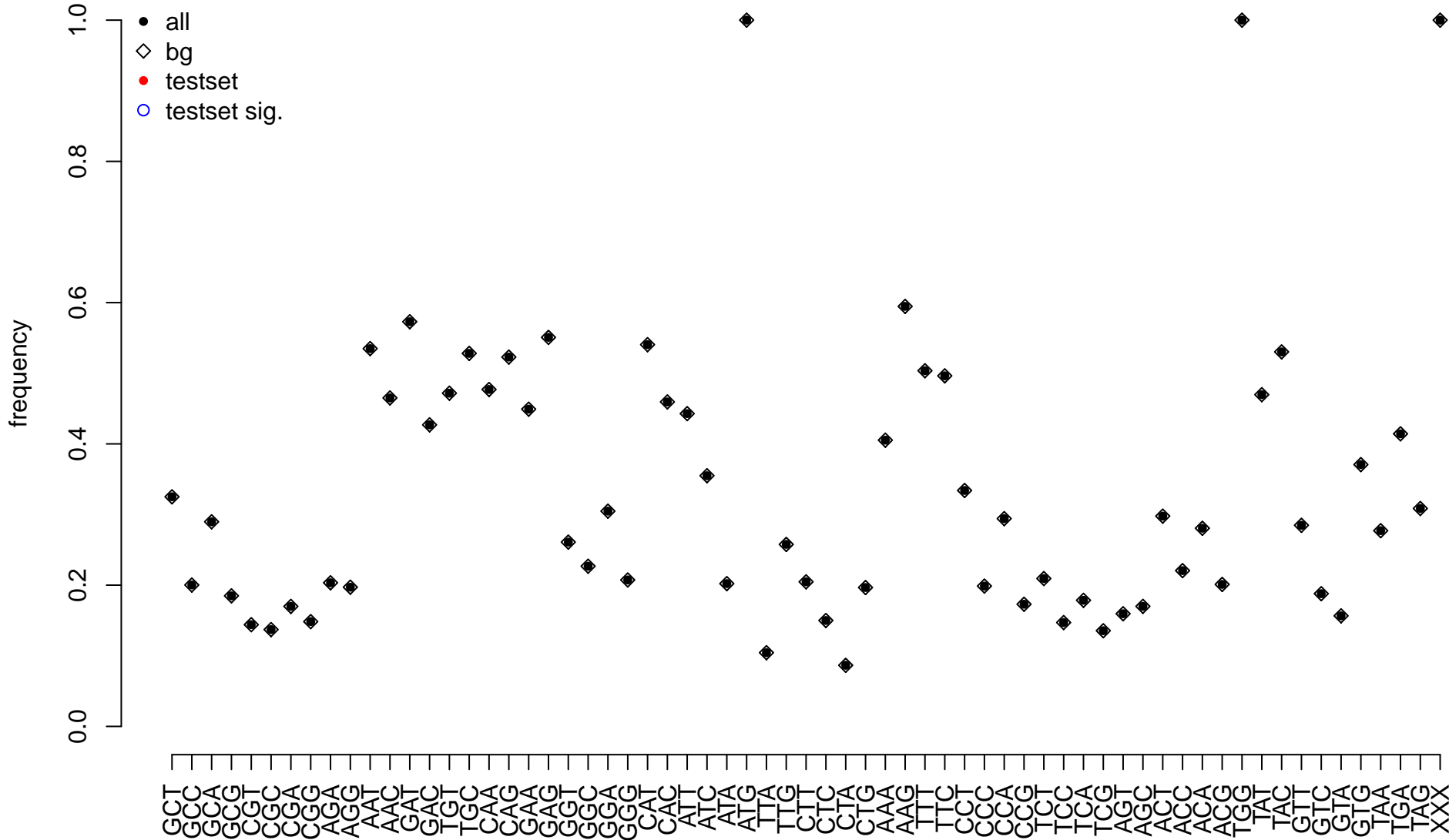
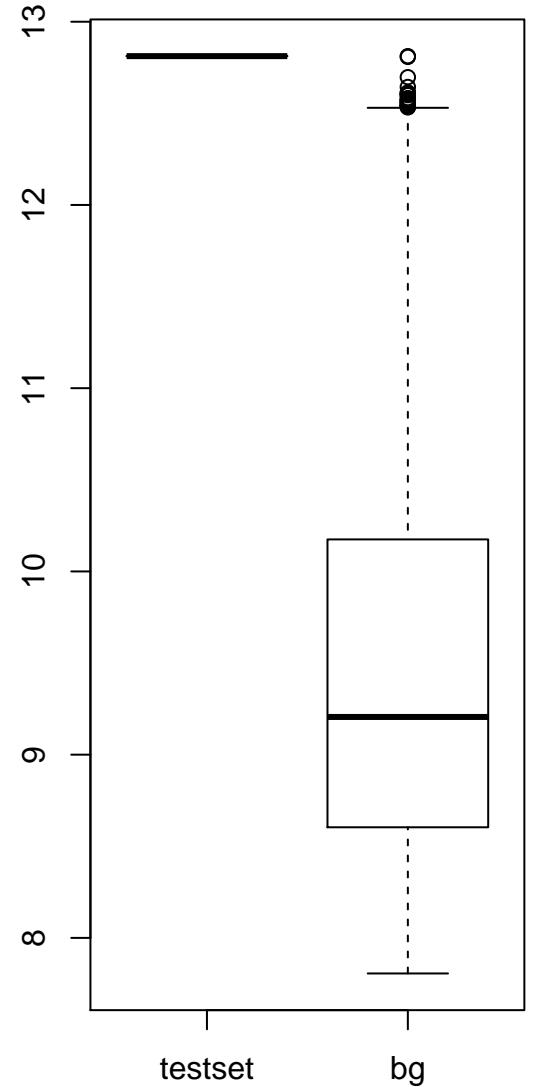


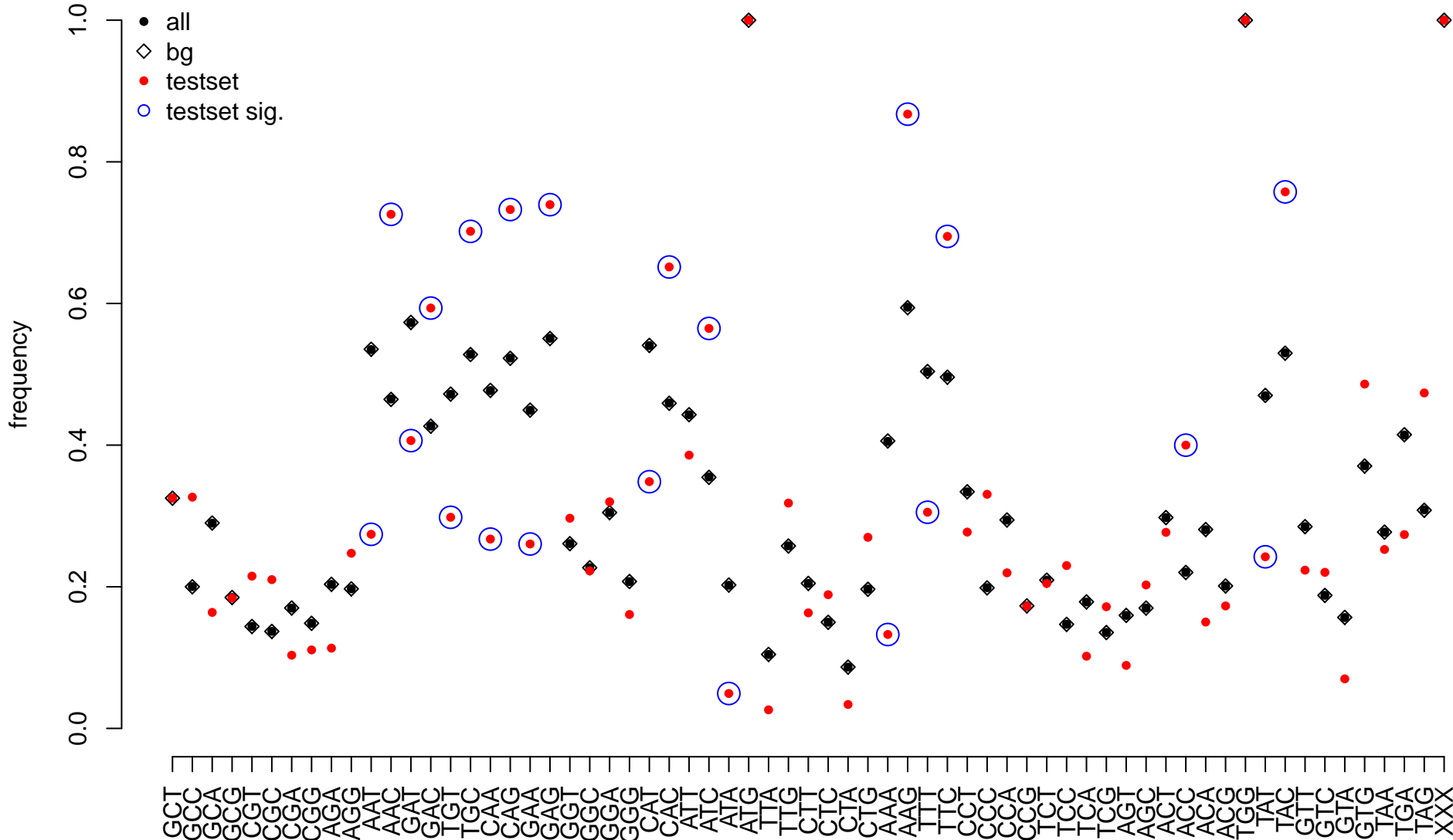
Codon usage frequency spectrum normalized per aa per gene  
group starting from highest expression values #genes: 0



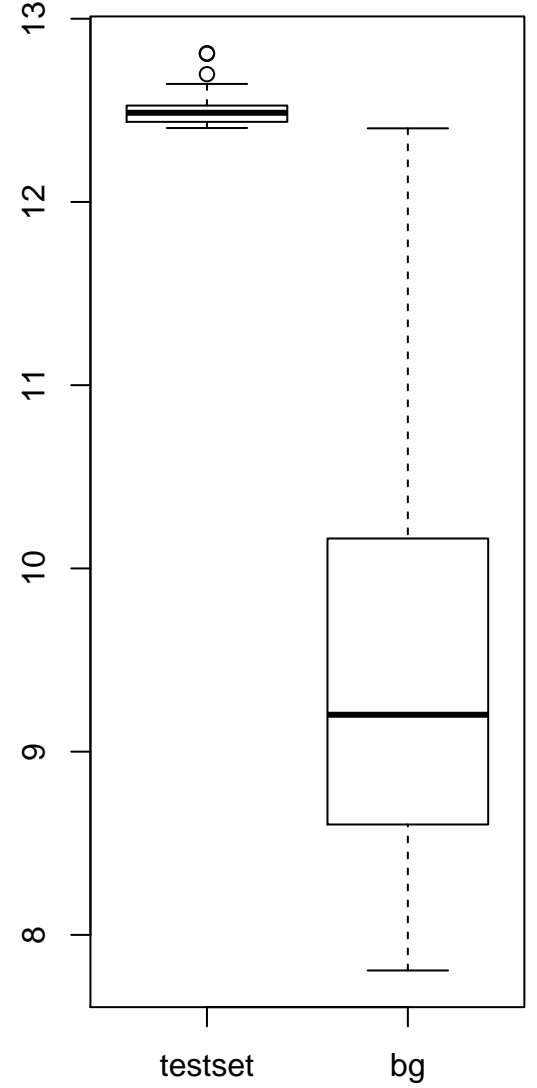
median expression values  
#genes: 0



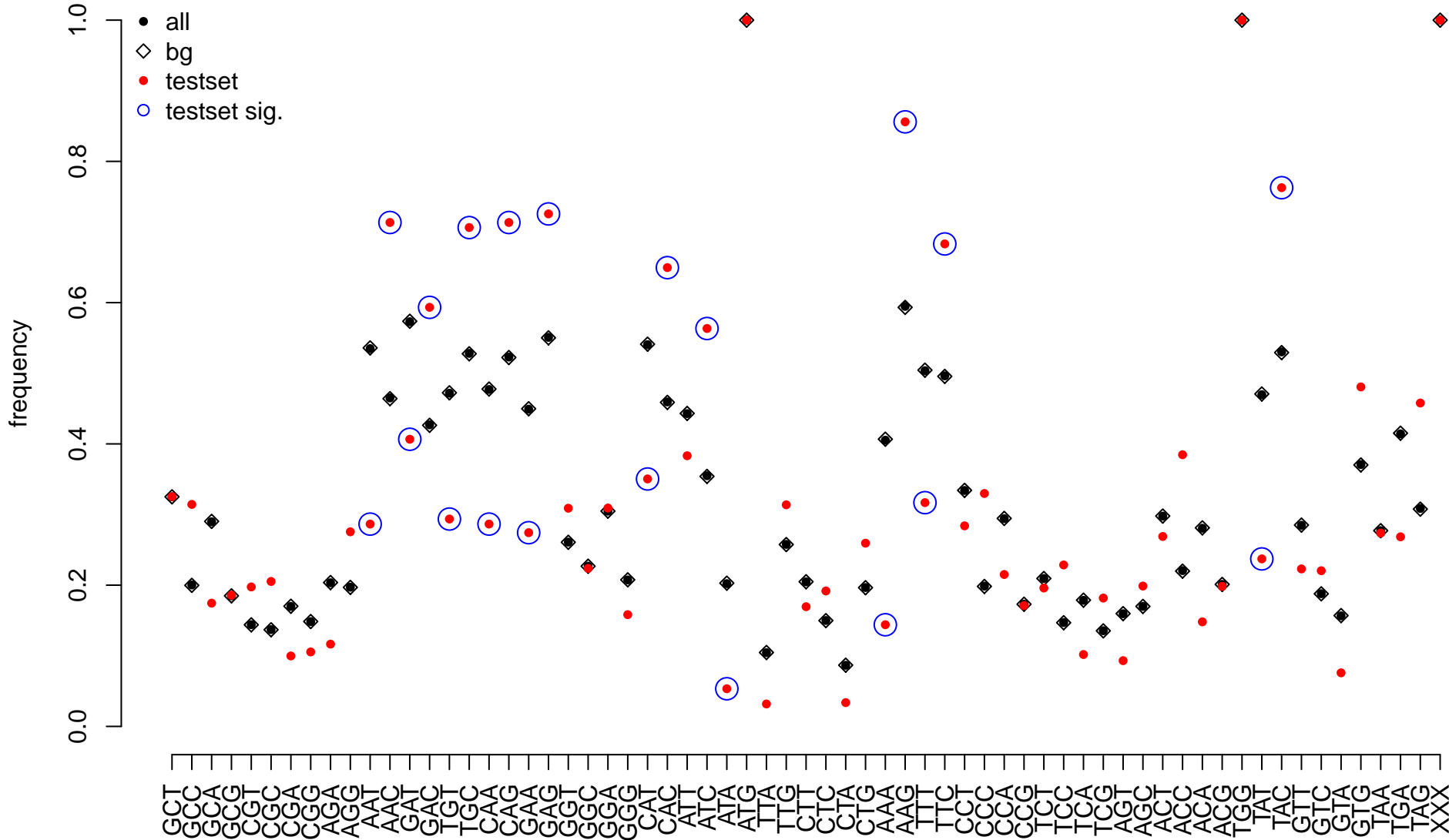
**Codon usage frequency spectrum normalized per aa per gene**  
**group starting from highest expression values #genes: 96**



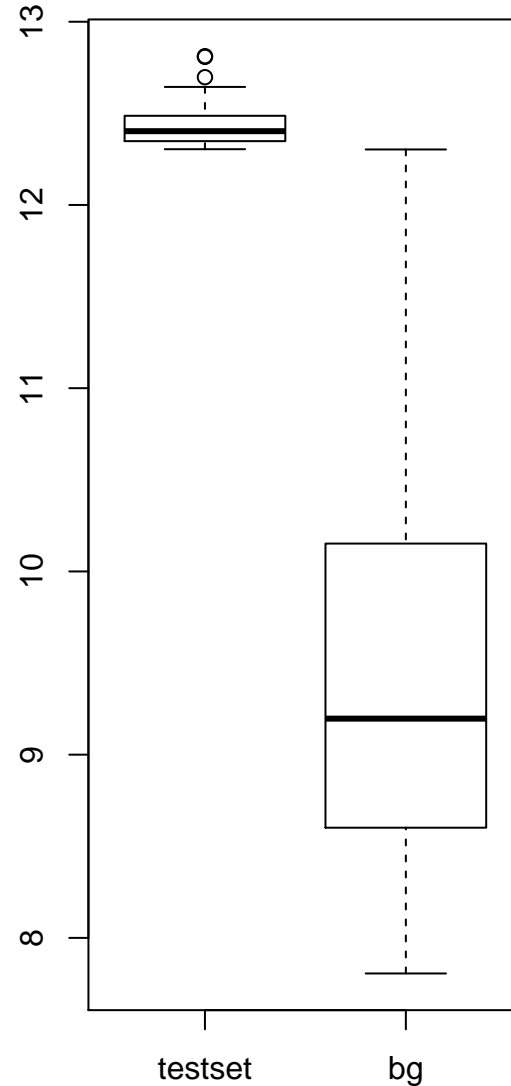
**median expression values**  
**#genes: 96**



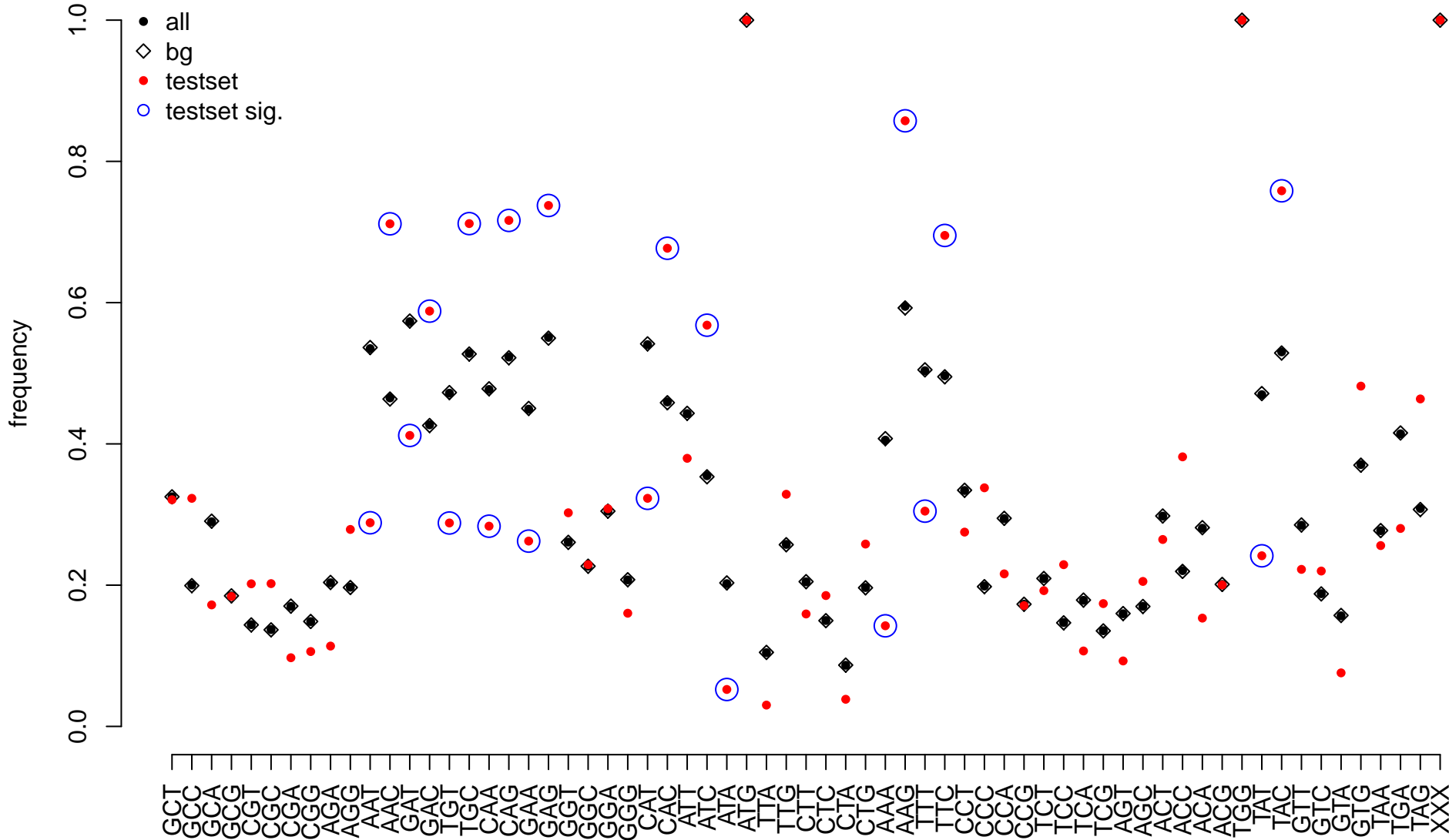
**Codon usage frequency spectrum normalized per aa per gene  
group starting from highest expression values #genes: 192**



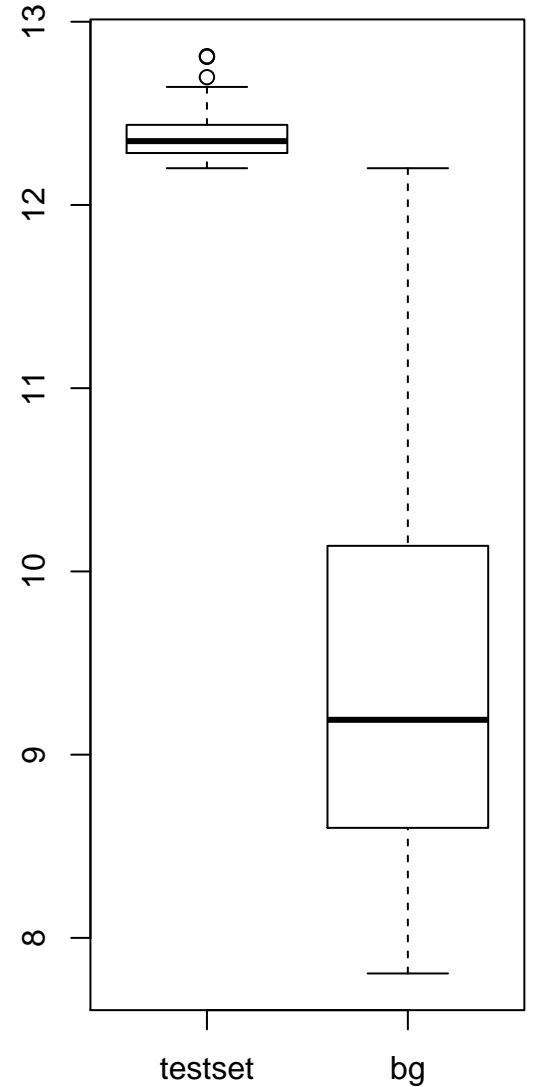
**median expression values  
#genes: 192**



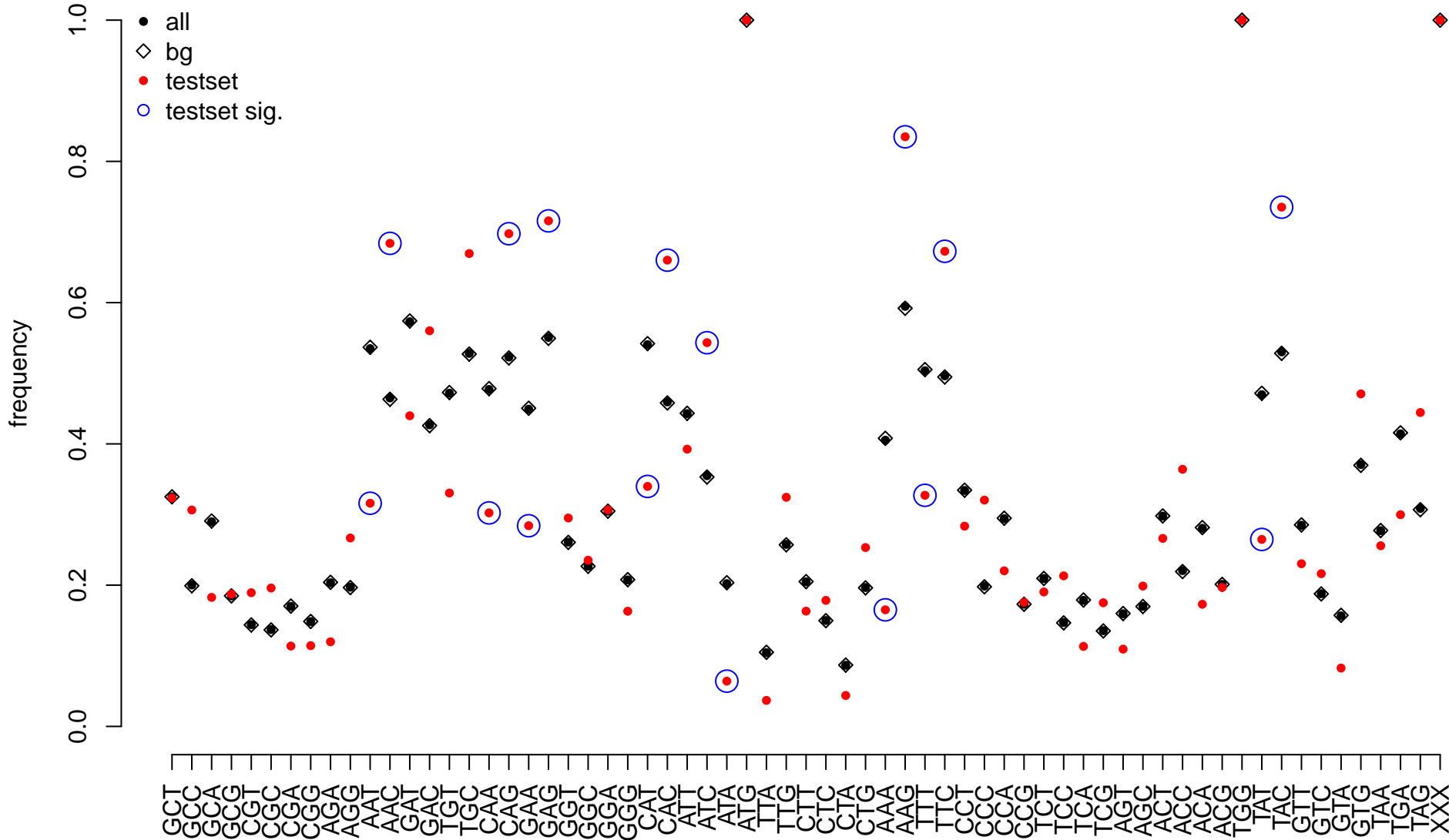
Codon usage frequency spectrum normalized per aa per gene  
group starting from highest expression values #genes: 291



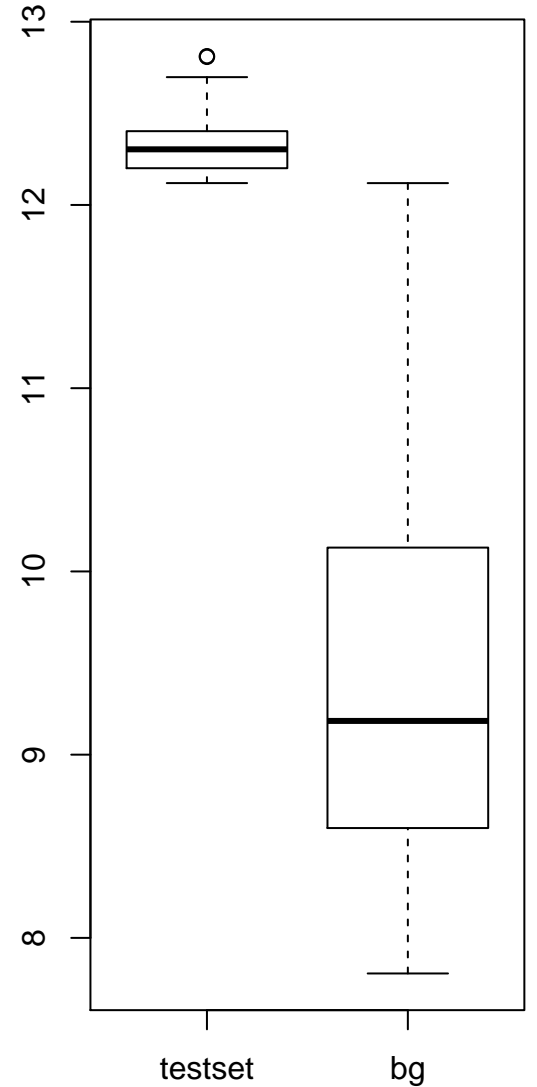
median expression values  
#genes: 291



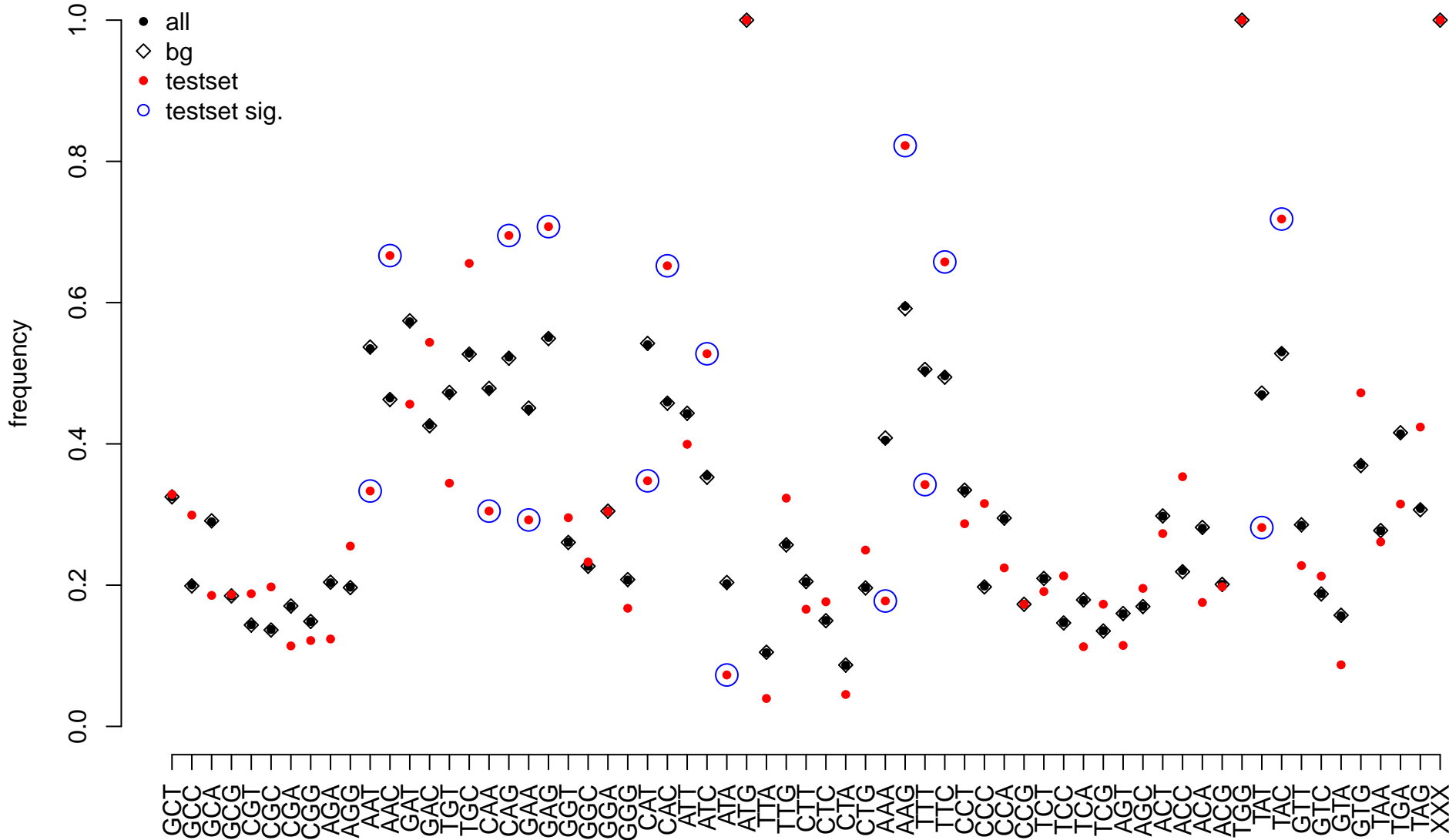
**Codon usage frequency spectrum normalized per aa per gene**  
**group starting from highest expression values #genes: 389**



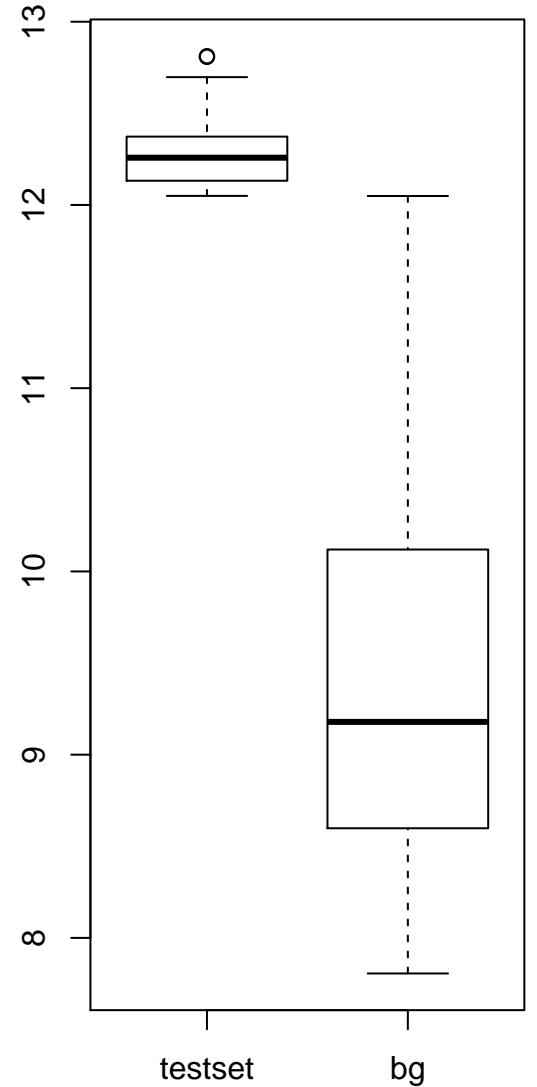
**median expression values**  
**#genes: 389**



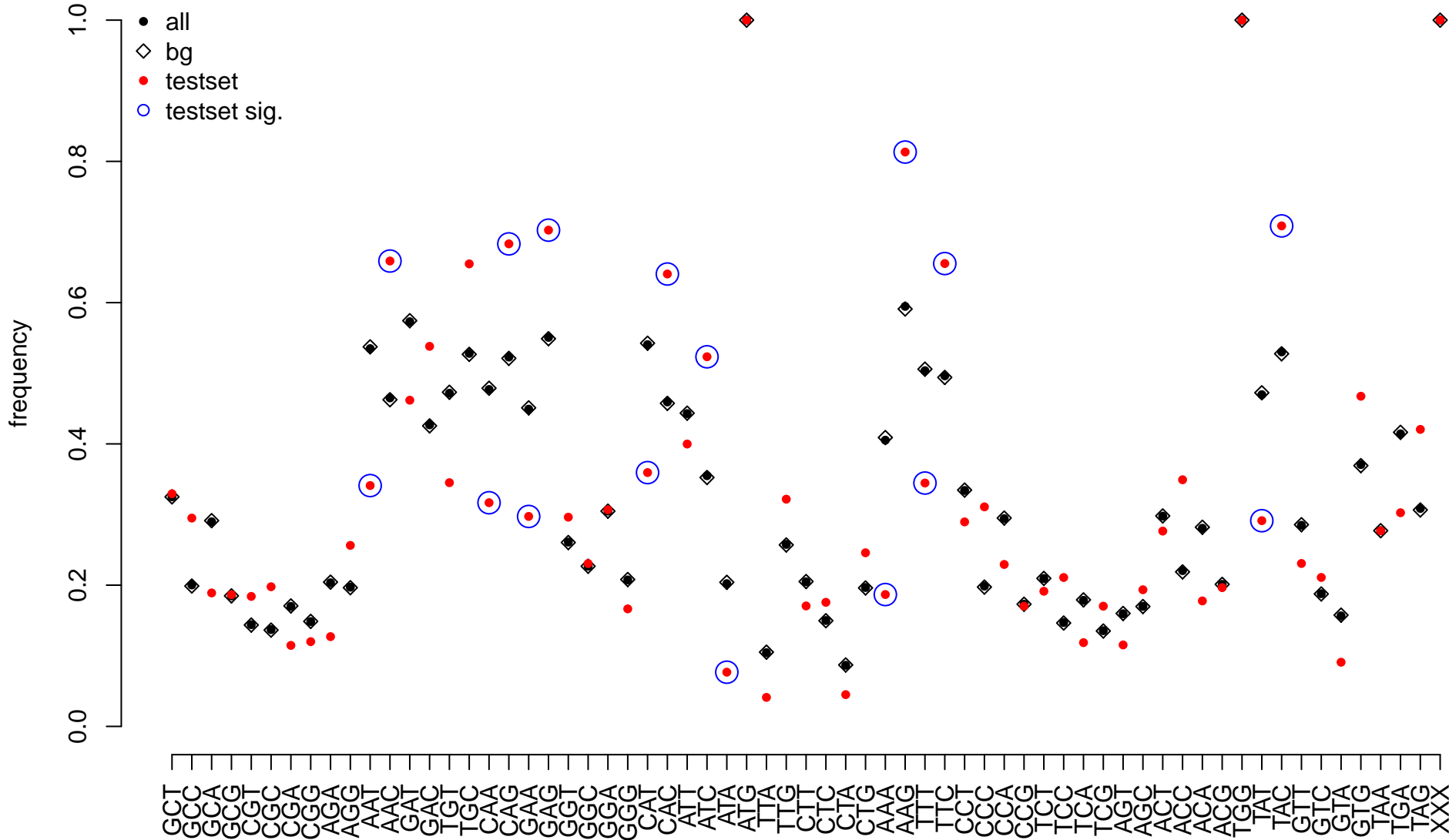
**Codon usage frequency spectrum normalized per aa per gene**  
**group starting from highest expression values #genes: 488**



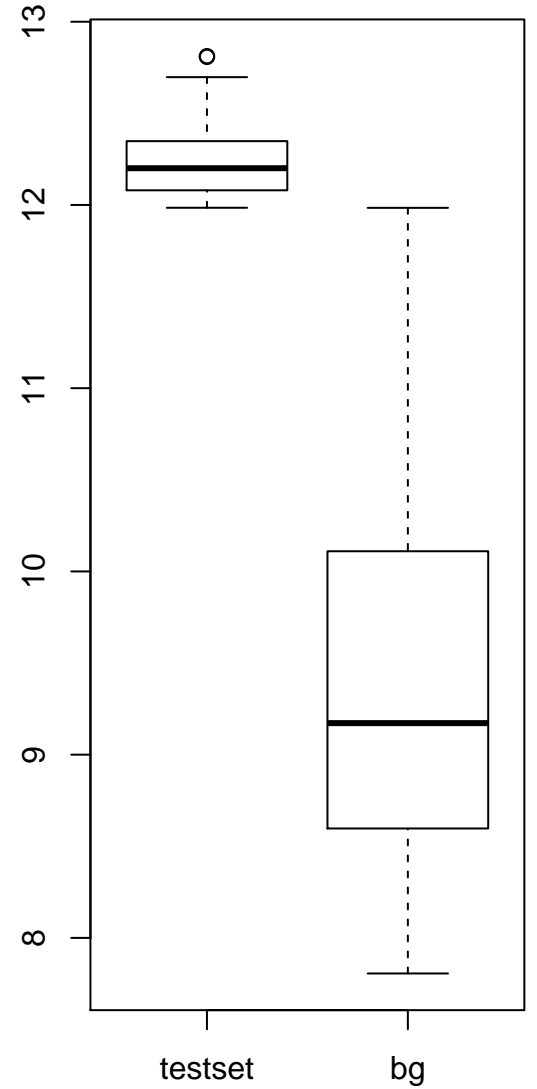
**median expression values**  
**#genes: 488**



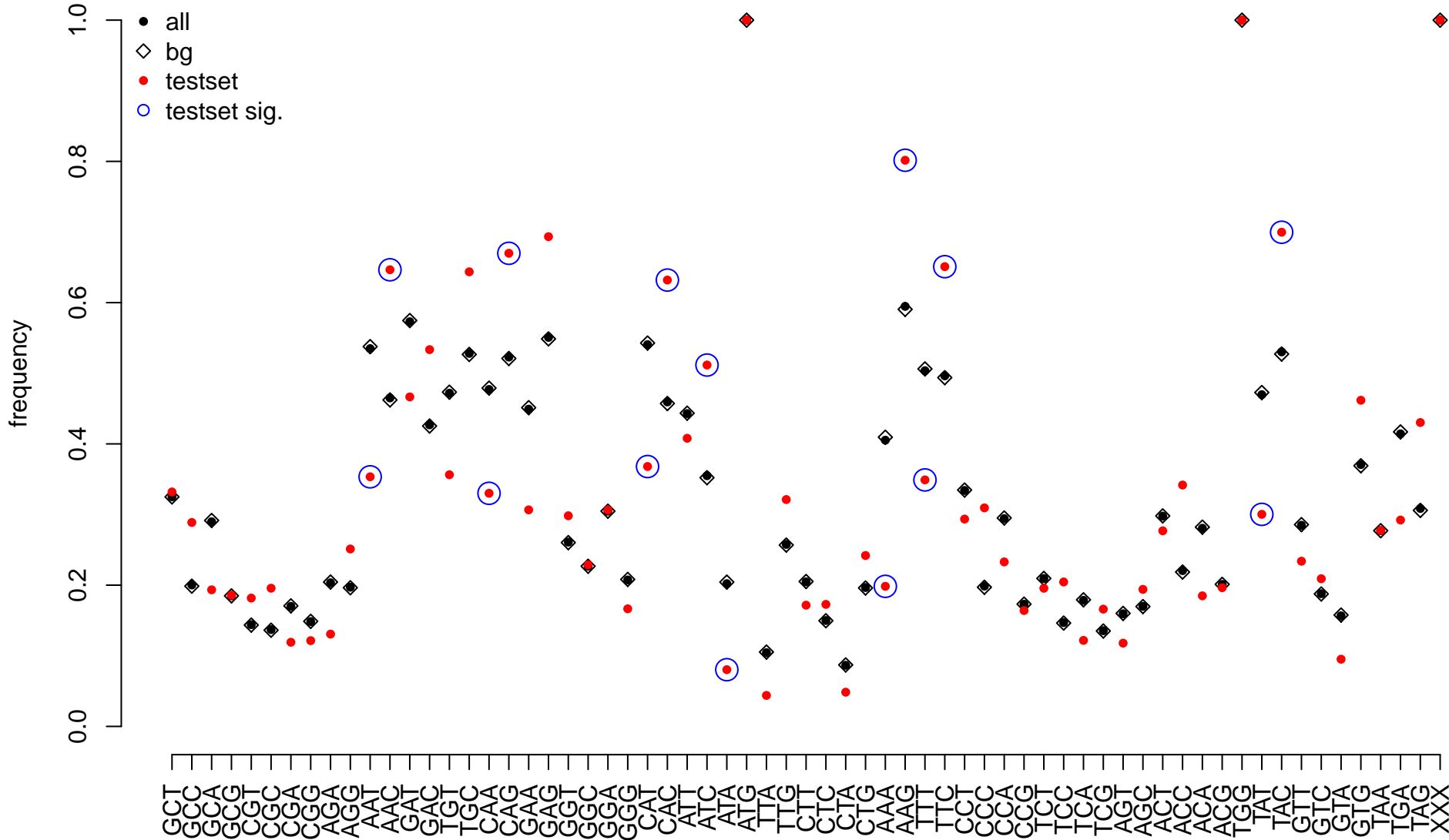
**Codon usage frequency spectrum normalized per aa per gene**  
**group starting from highest expression values #genes: 587**



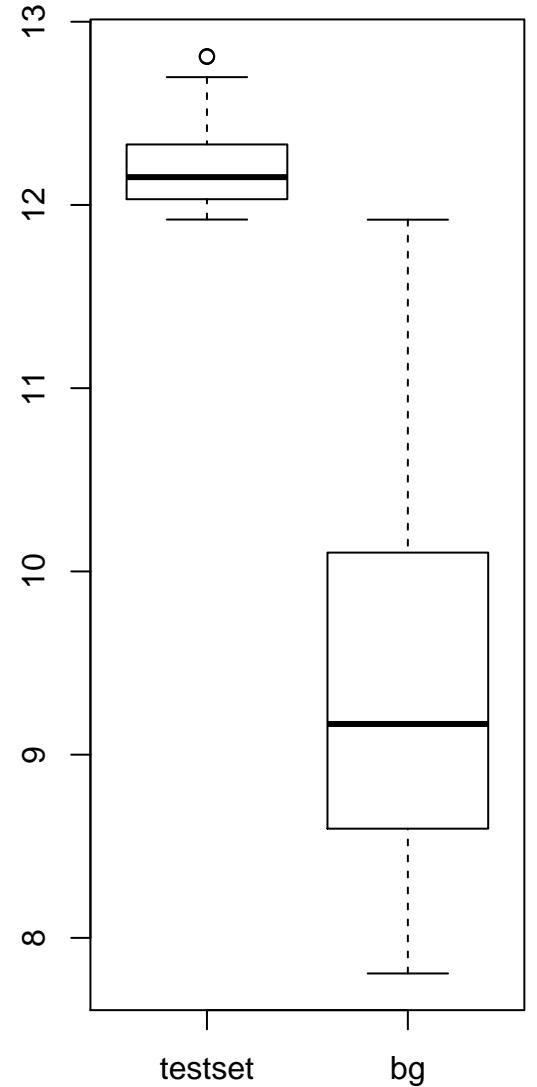
**median expression values**  
**#genes: 587**



Codon usage frequency spectrum normalized per aa per gene  
group starting from highest expression values #genes: 683



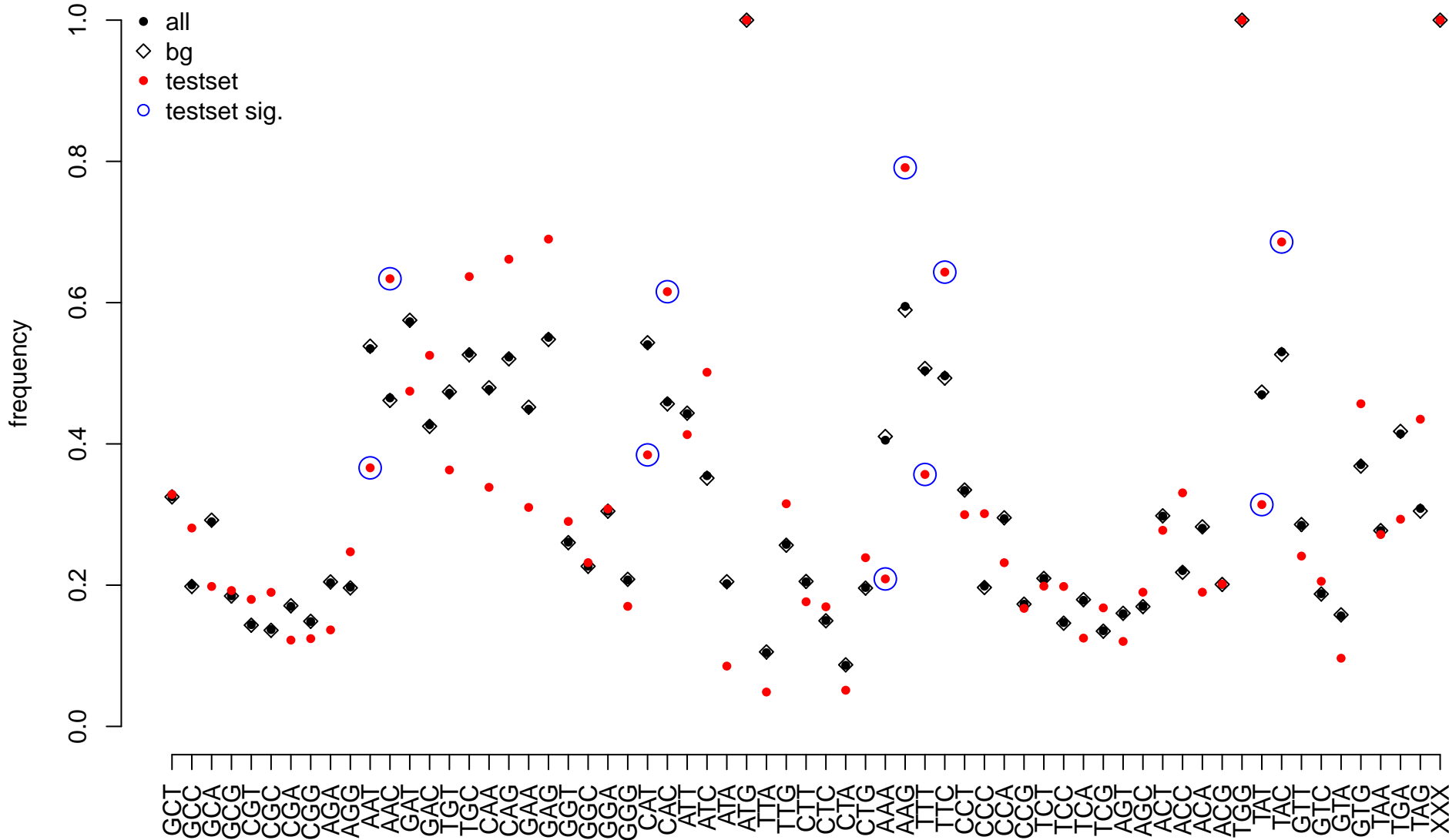
median expression values  
#genes: 683



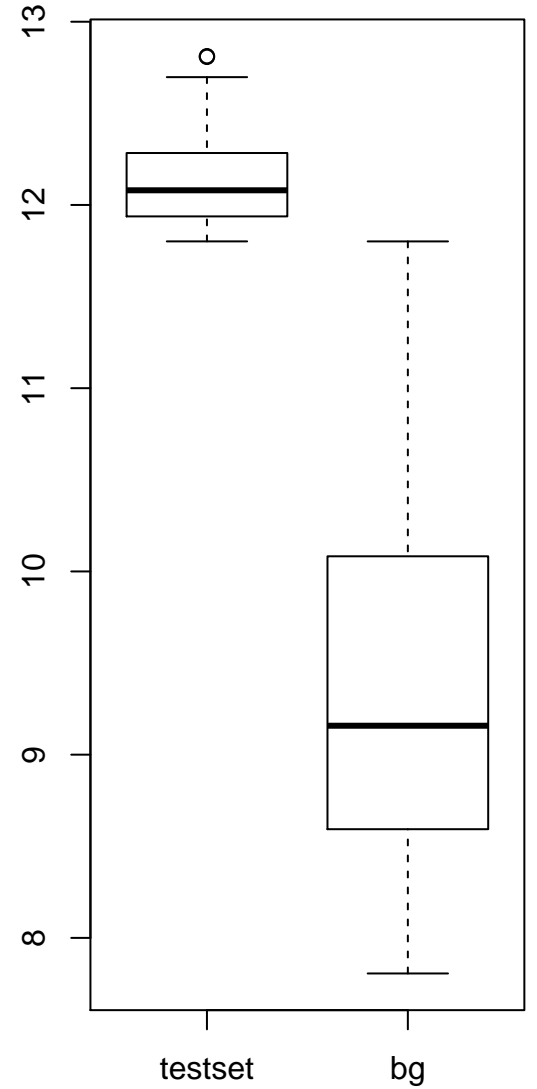




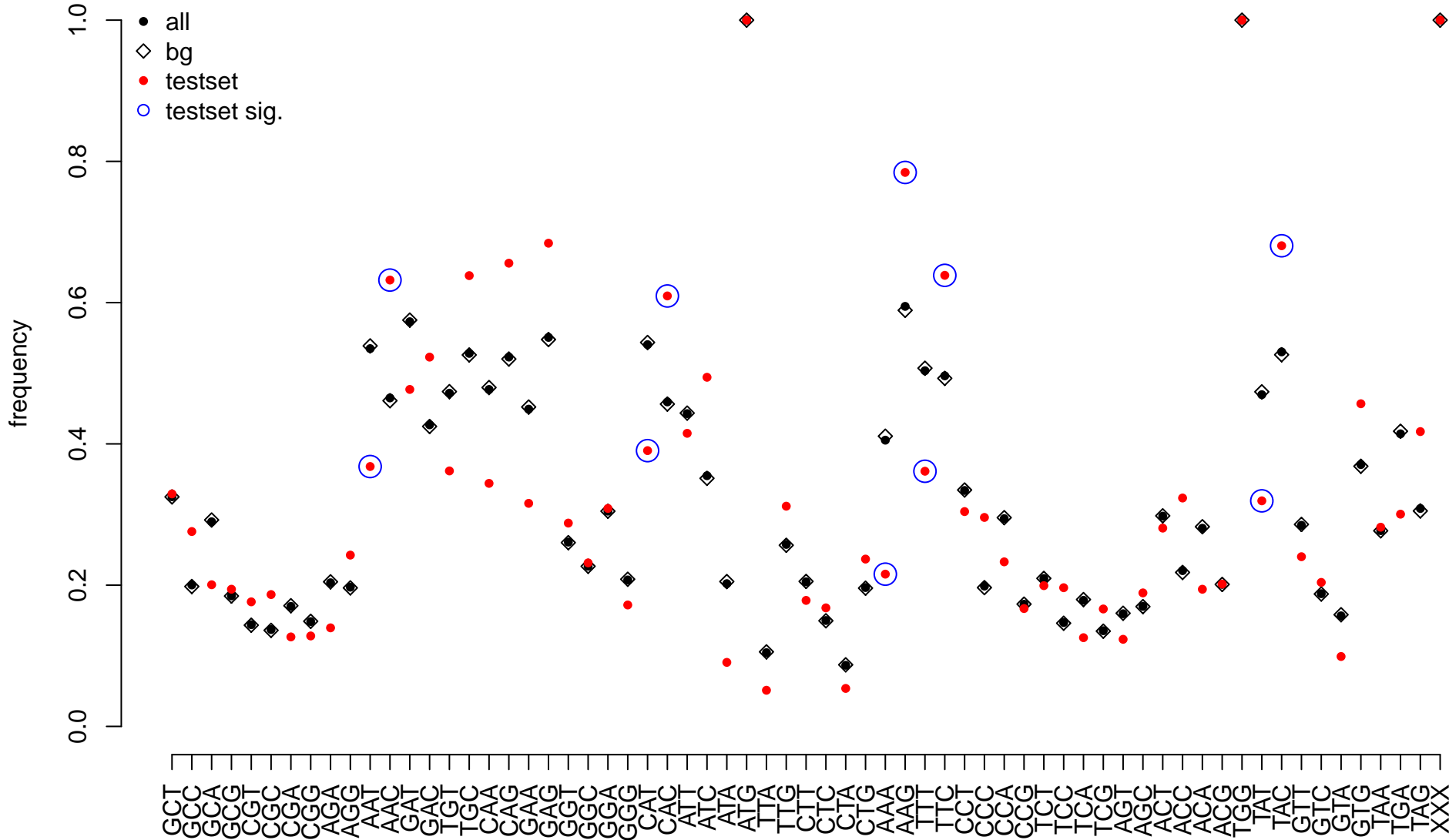
**Codon usage frequency spectrum normalized per aa per gene**  
**group starting from highest expression values #genes: 880**



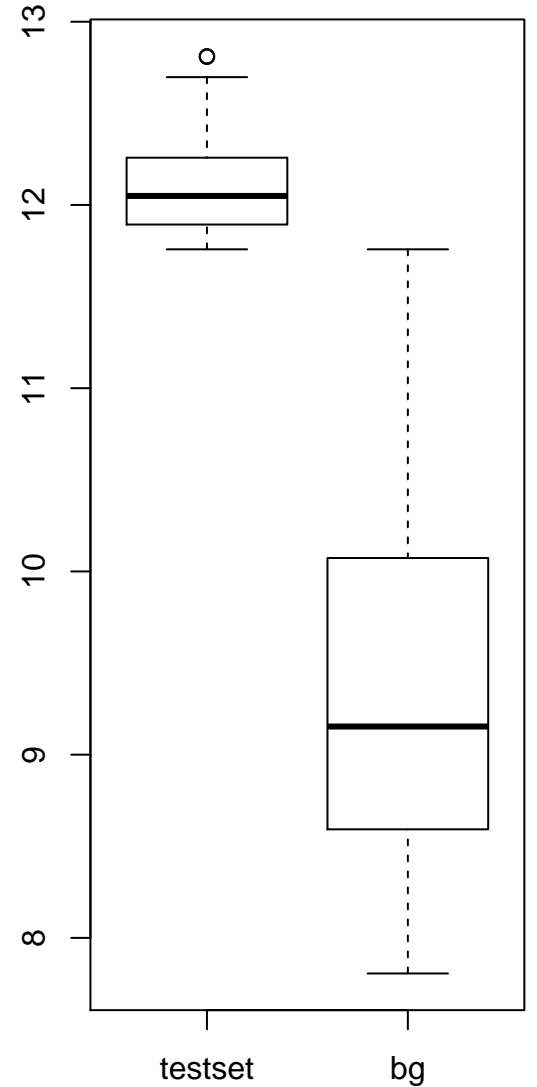
**median expression values**  
**#genes: 880**



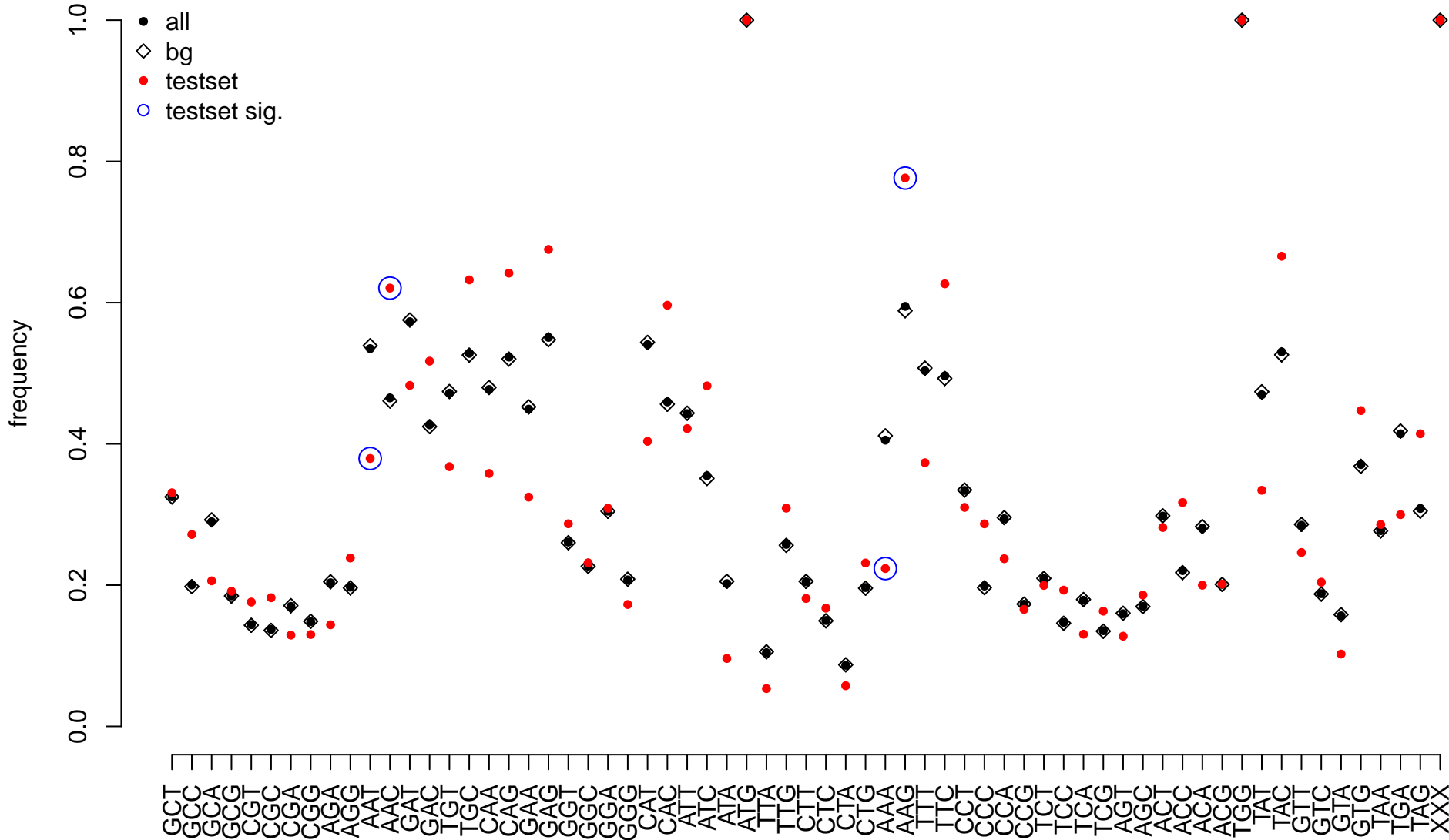
**Codon usage frequency spectrum normalized per aa per gene**  
**group starting from highest expression values #genes: 980**



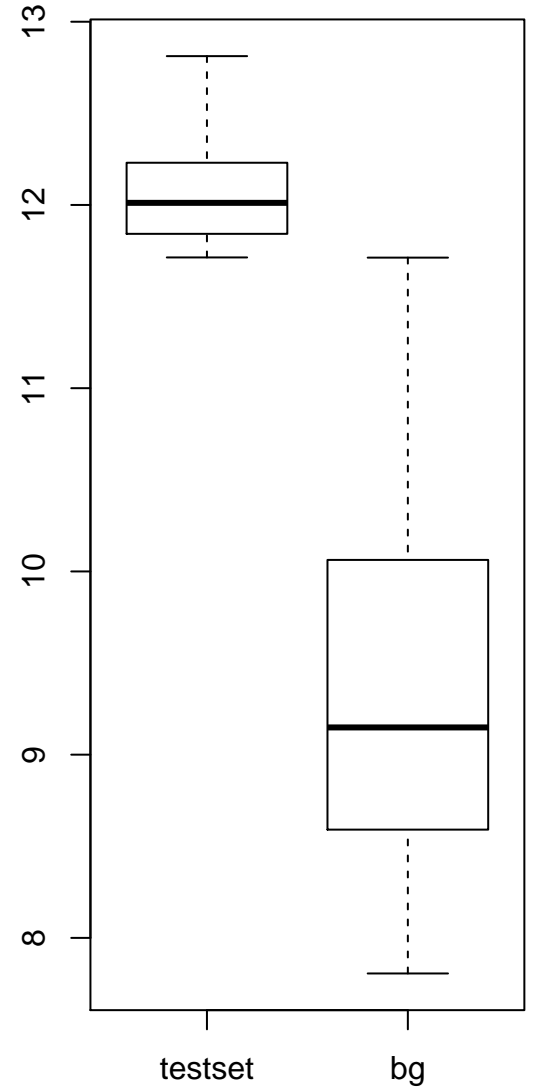
**median expression values**  
**#genes: 980**



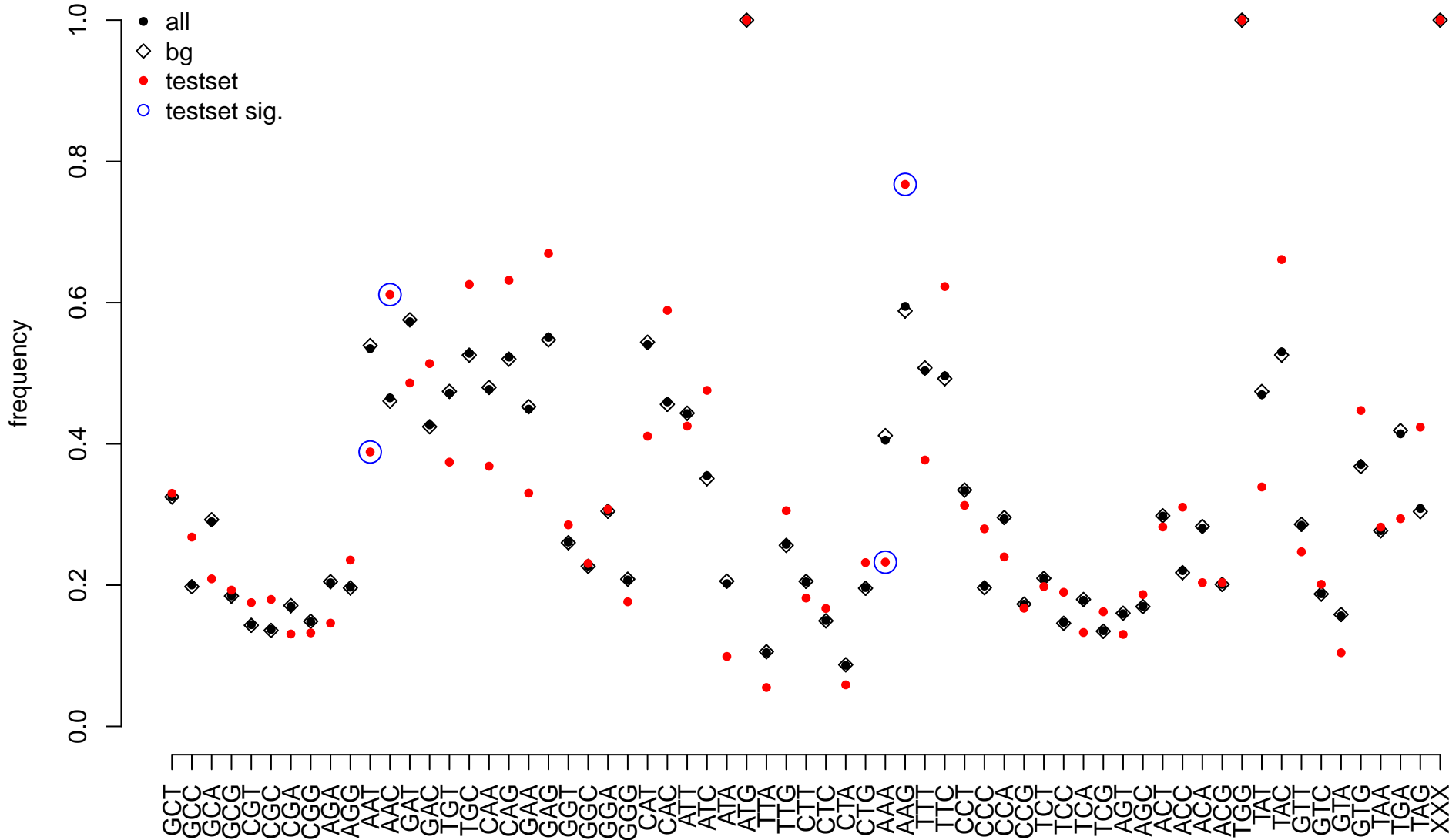
Codon usage frequency spectrum normalized per aa per gene  
group starting from highest expression values #genes: 1080



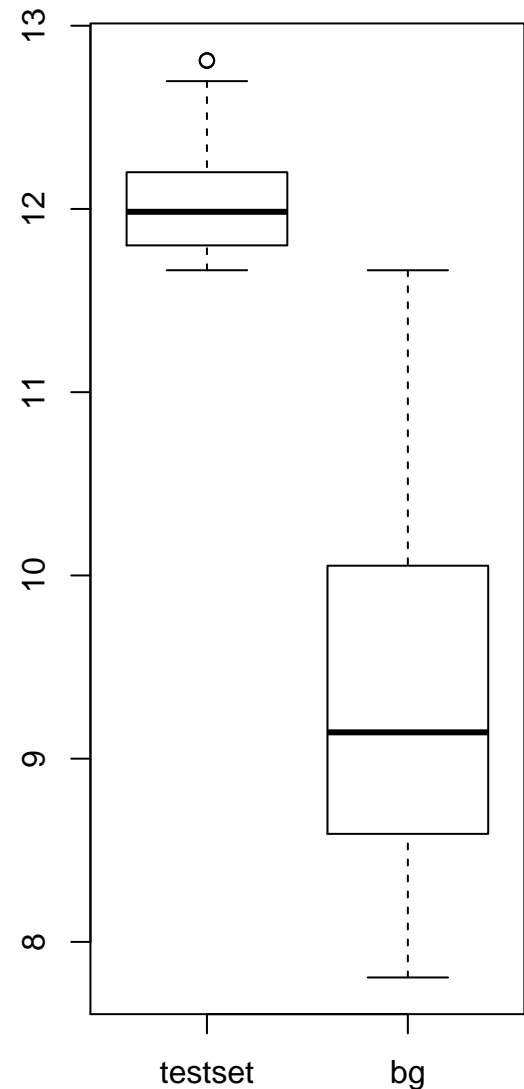
median expression values  
#genes: 1080



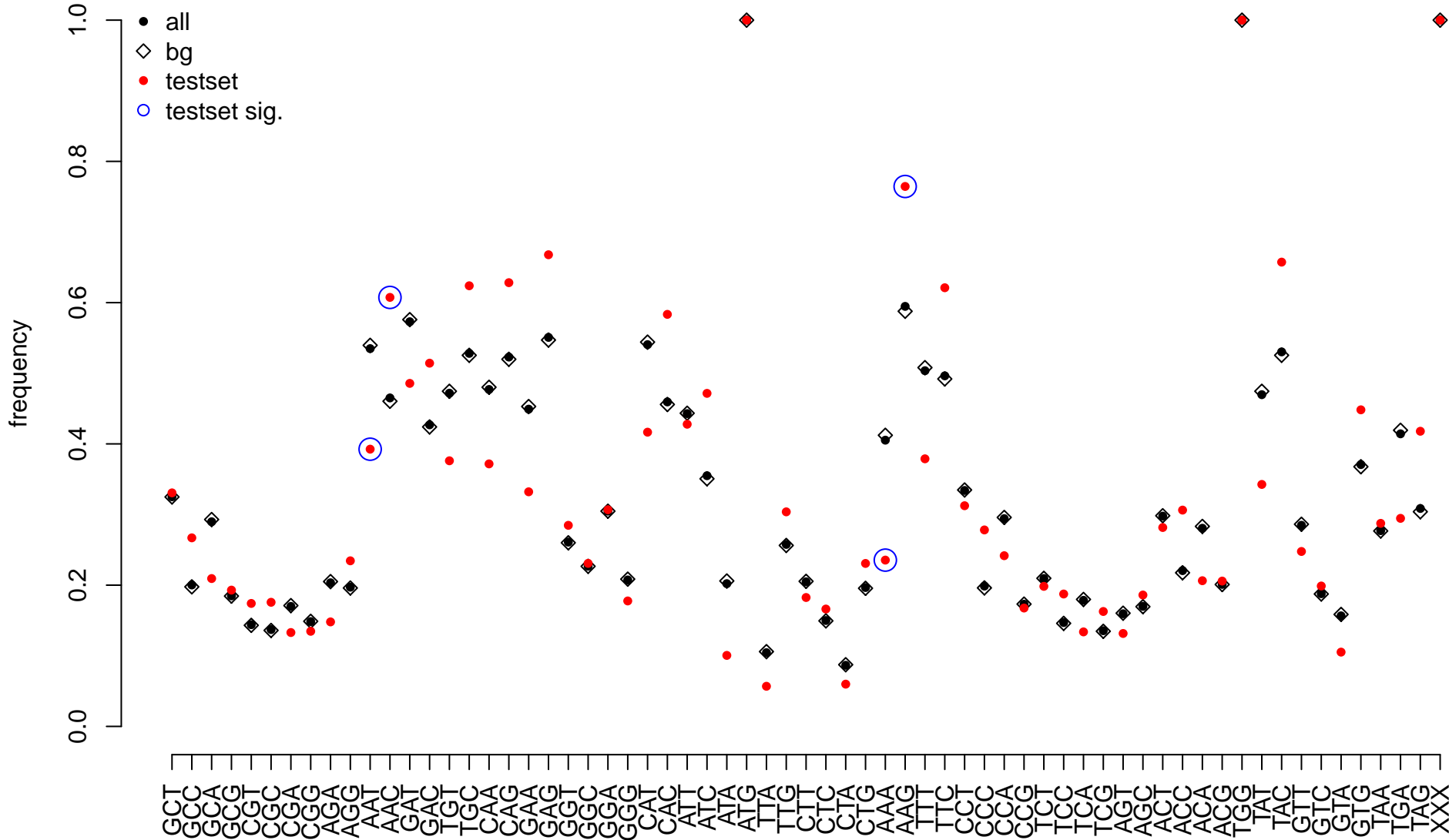
Codon usage frequency spectrum normalized per aa per gene  
group starting from highest expression values #genes: 1179



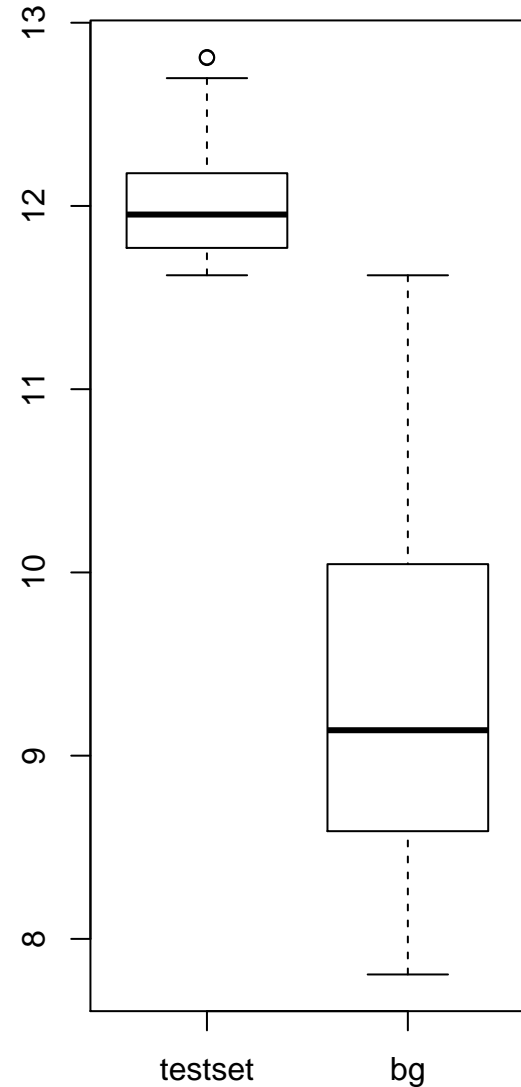
median expression values  
#genes: 1179



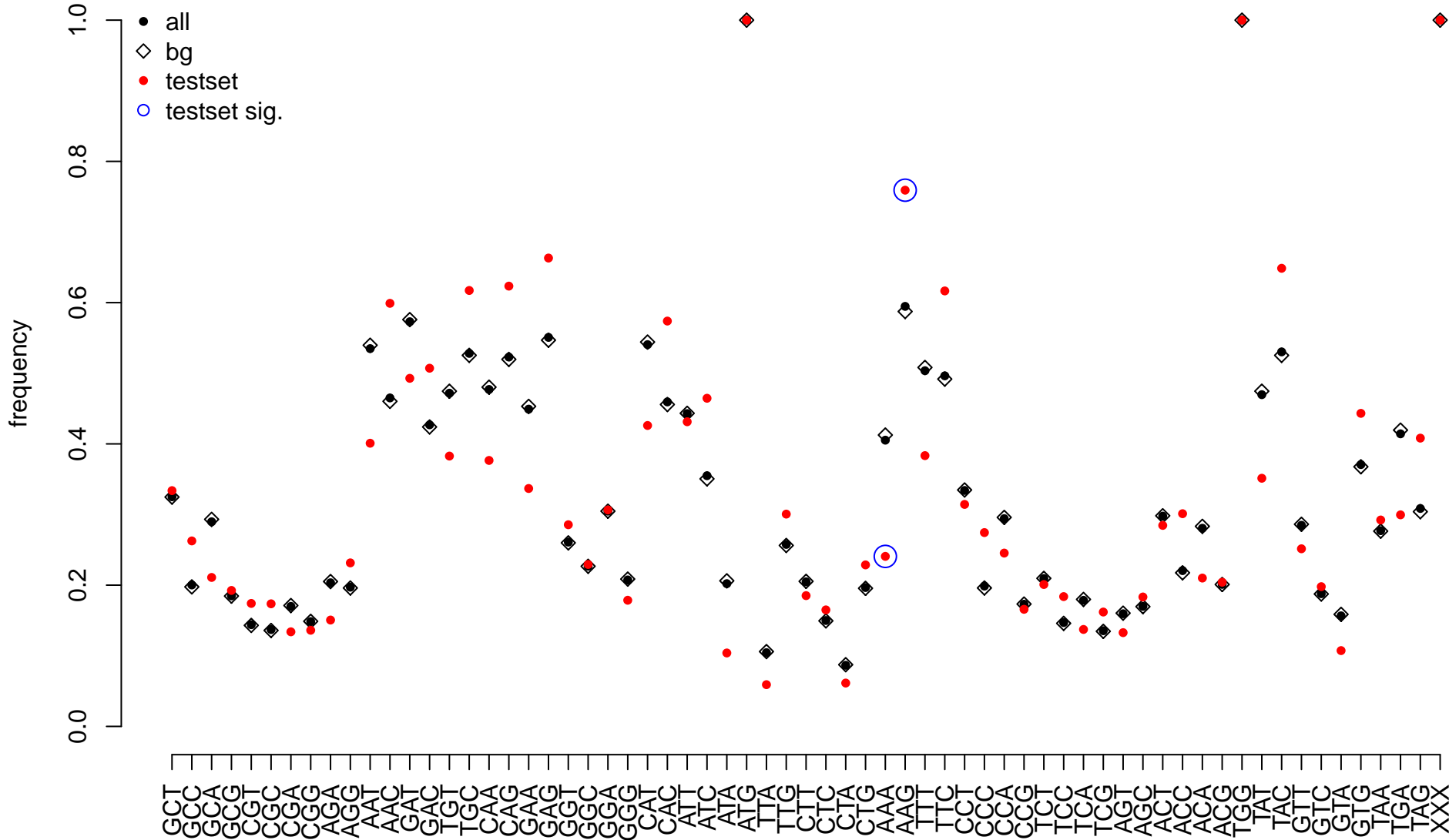
Codon usage frequency spectrum normalized per aa per gene  
group starting from highest expression values #genes: 1279



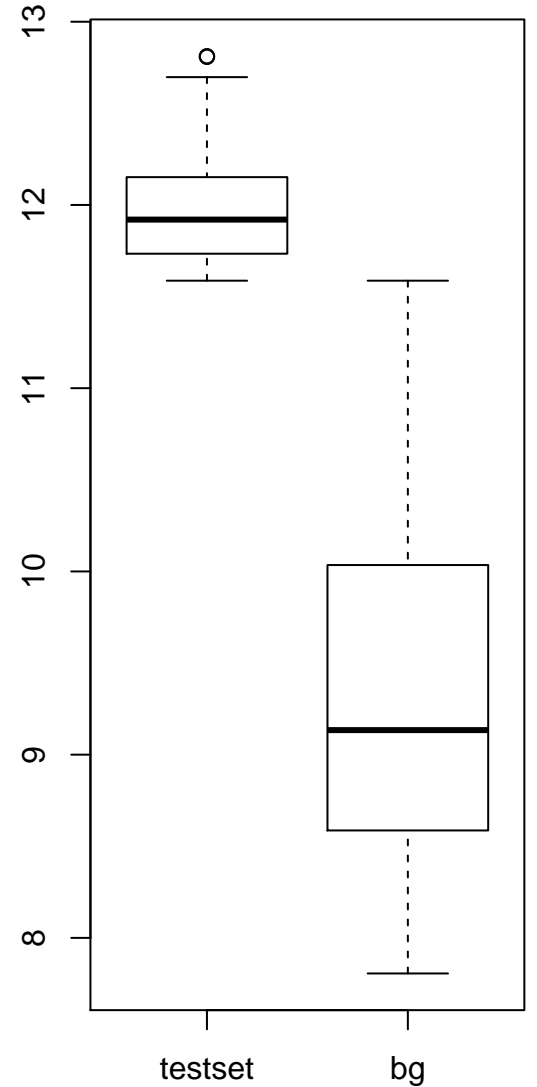
median expression values  
#genes: 1279



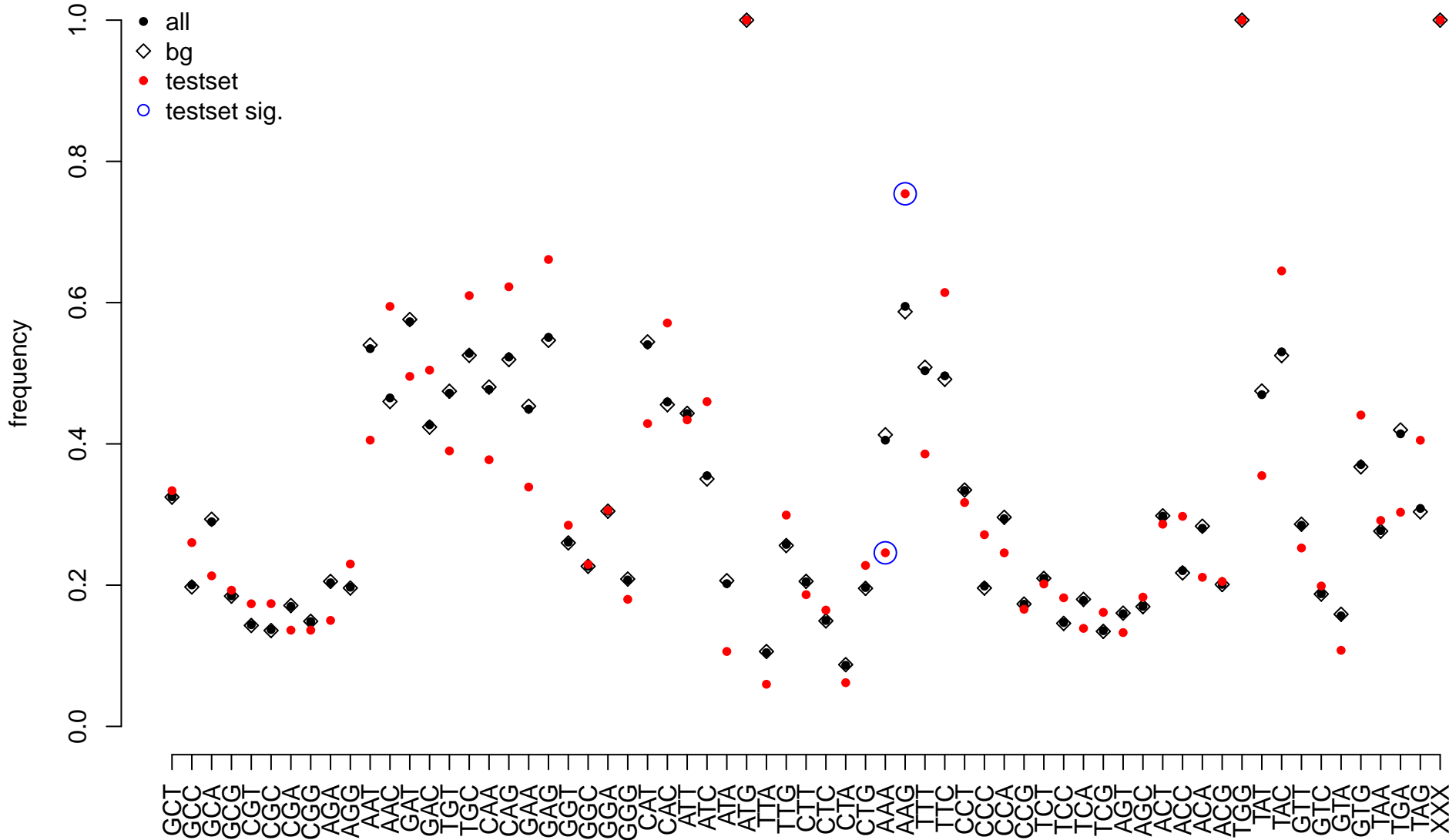
Codon usage frequency spectrum normalized per aa per gene  
group starting from highest expression values #genes: 1378



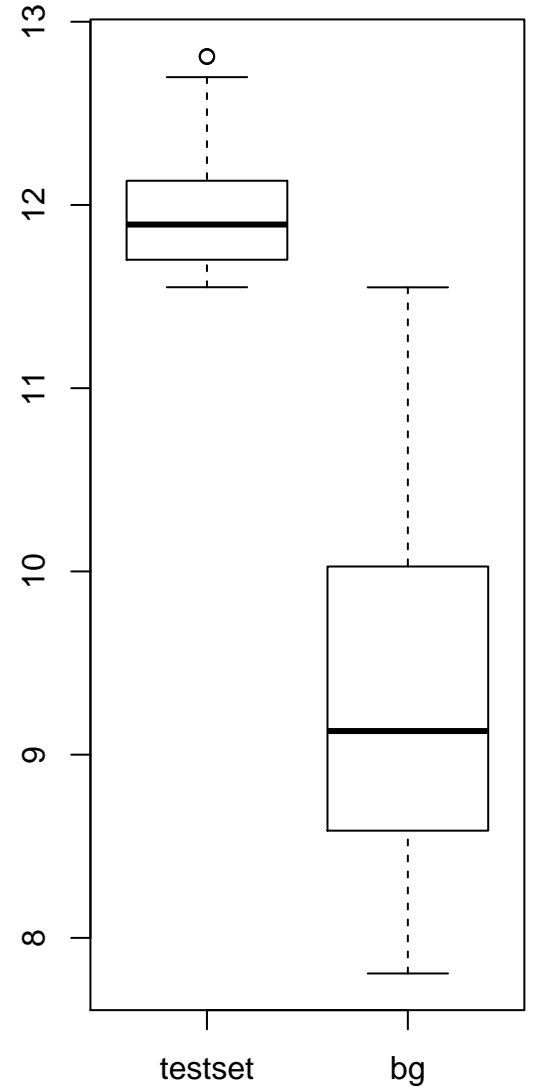
median expression values  
#genes: 1378



Codon usage frequency spectrum normalized per aa per gene  
group starting from highest expression values #genes: 1477

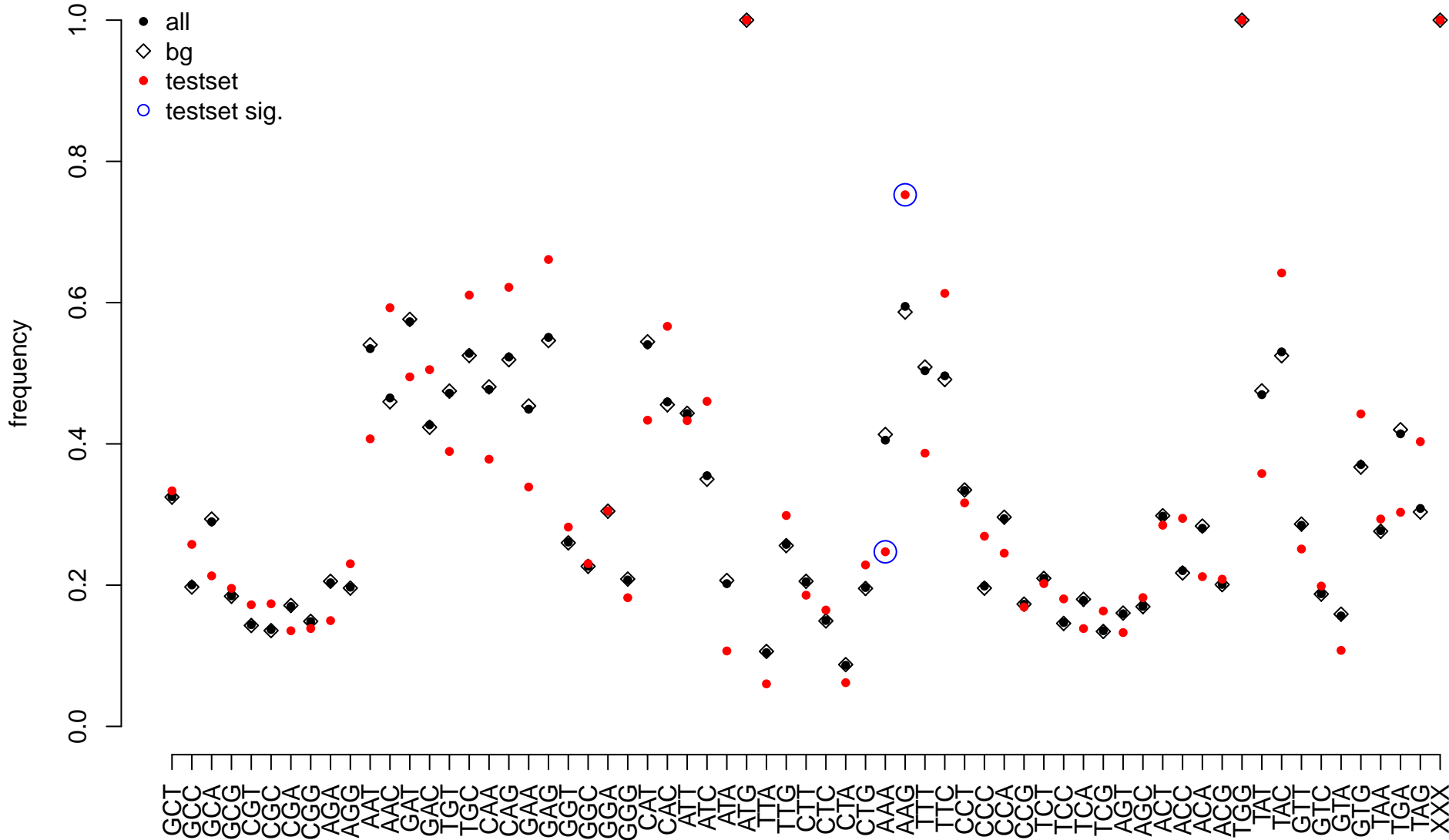


median expression values  
#genes: 1477

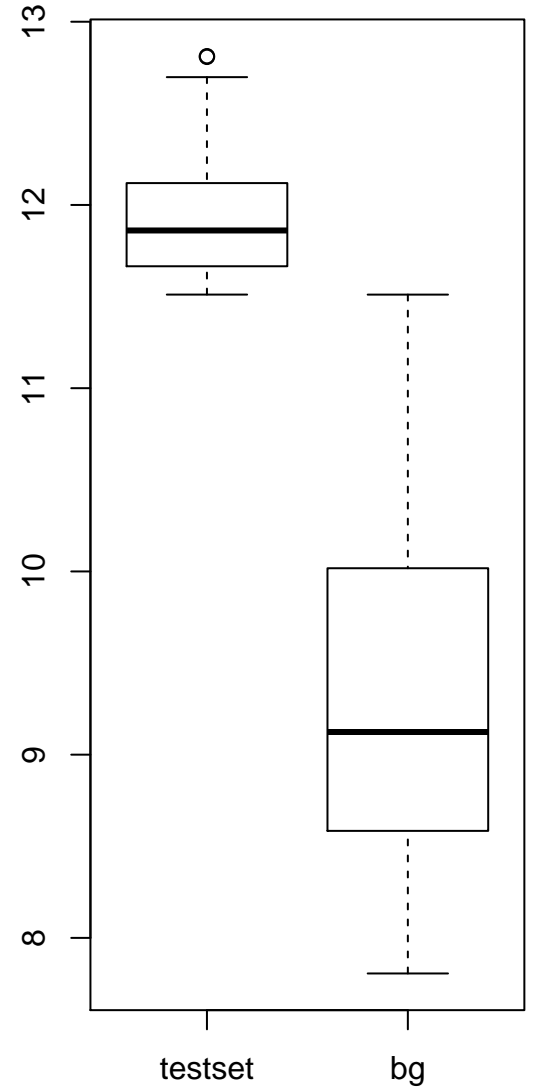




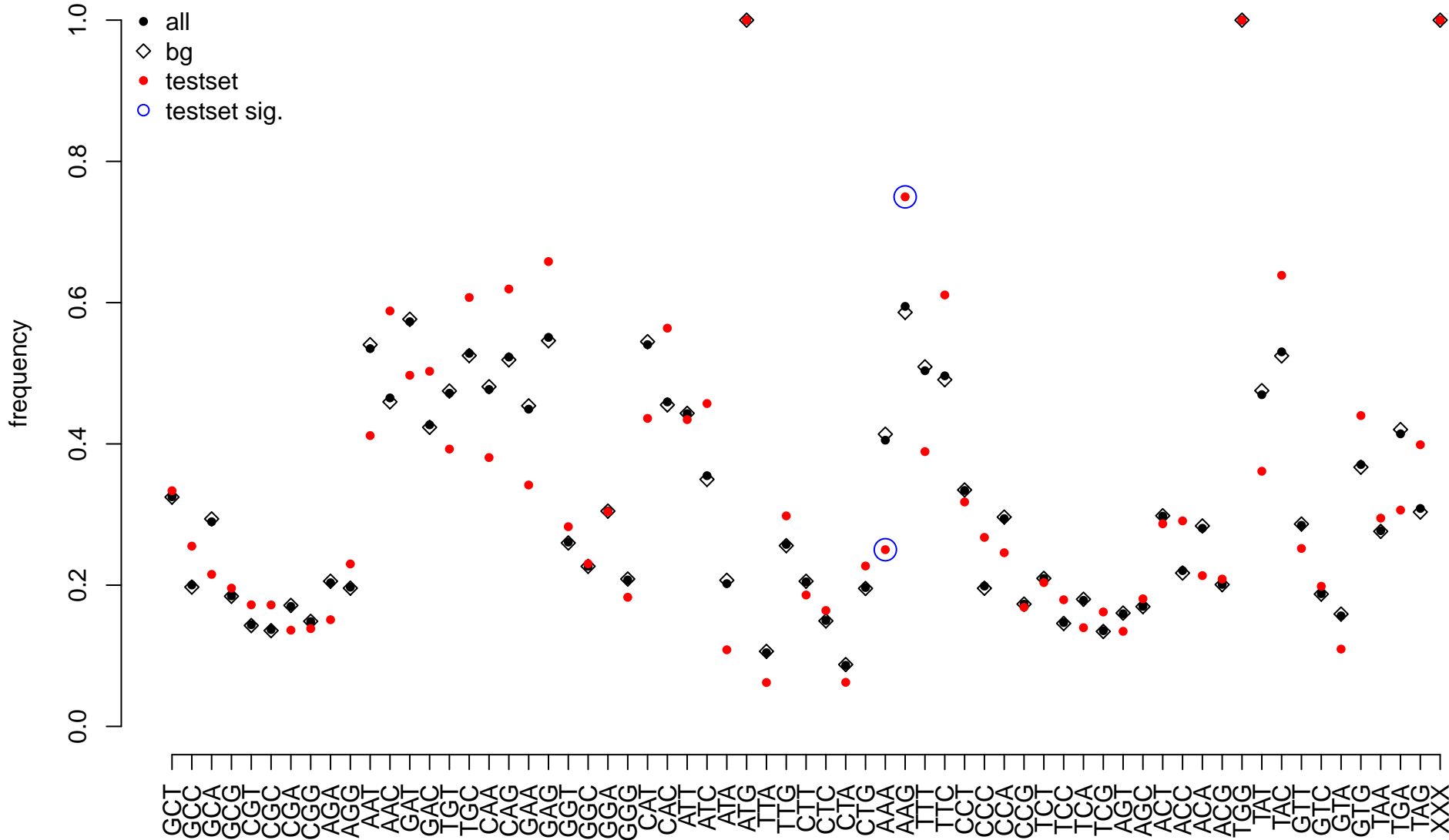
Codon usage frequency spectrum normalized per aa per gene  
group starting from highest expression values #genes: 1576



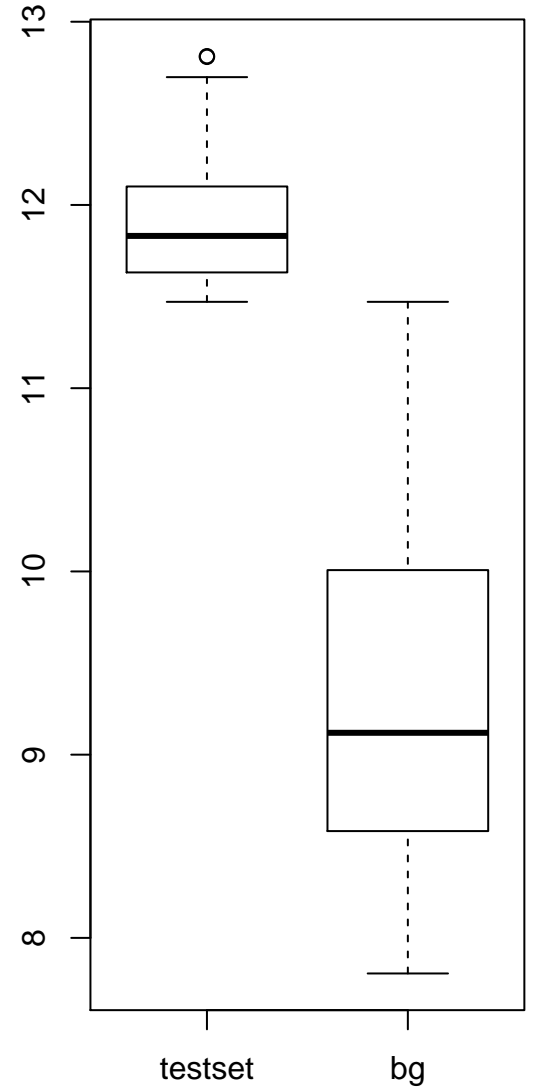
median expression values  
#genes: 1576



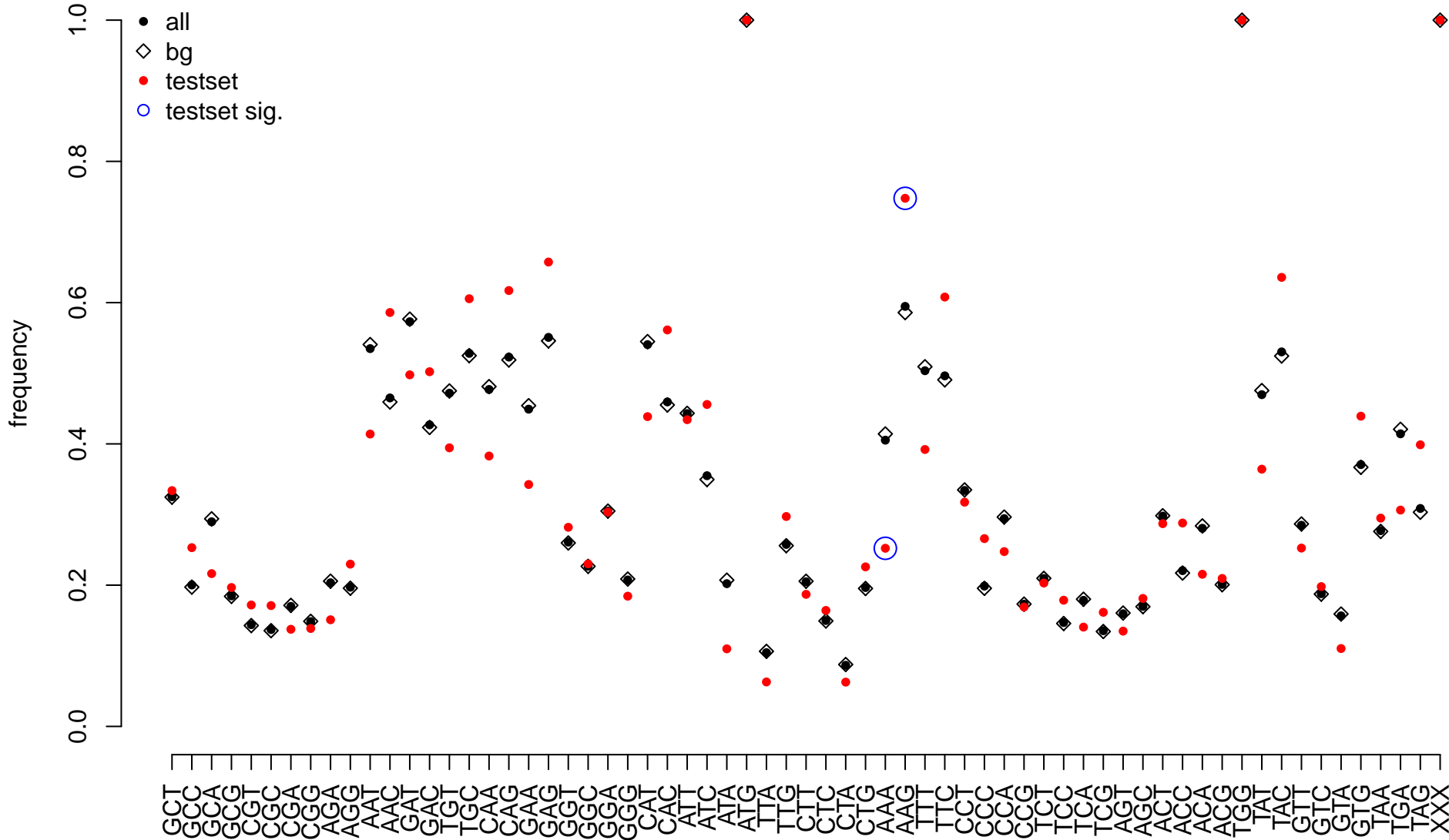
**Codon usage frequency spectrum normalized per aa per gene  
group starting from highest expression values #genes: 1672**



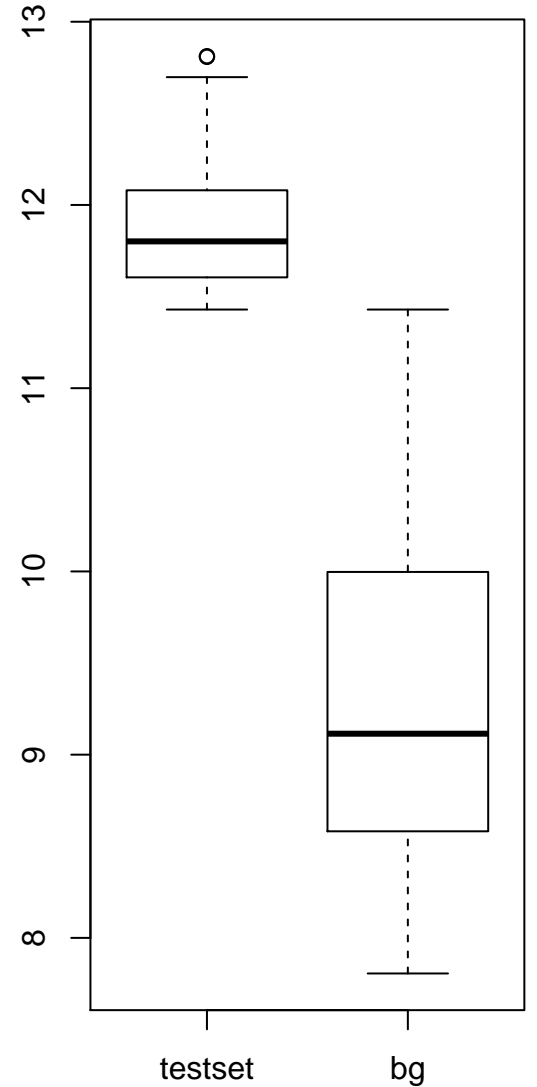
**median expression values  
#genes: 1672**



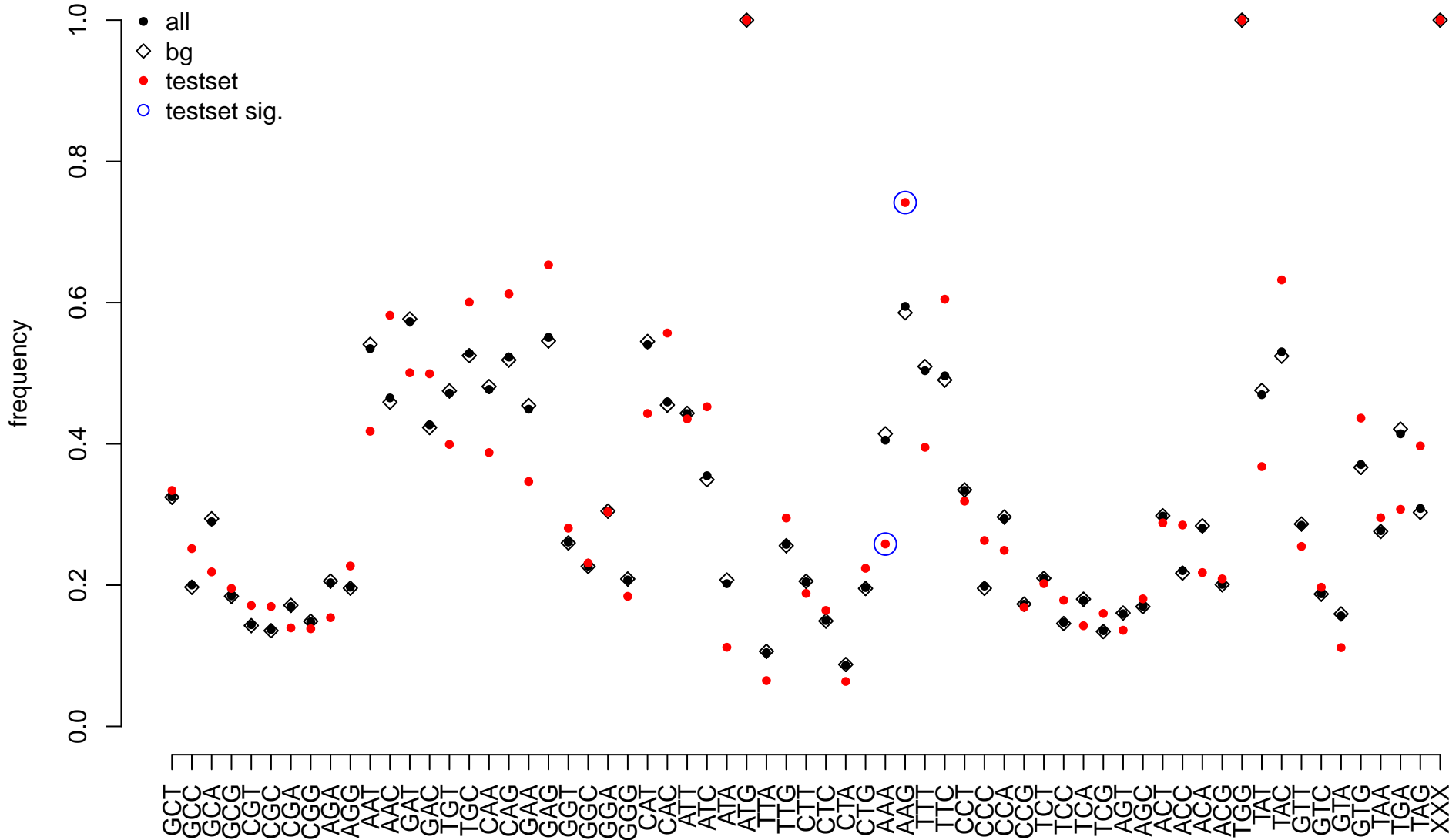
Codon usage frequency spectrum normalized per aa per gene  
group starting from highest expression values #genes: 1771



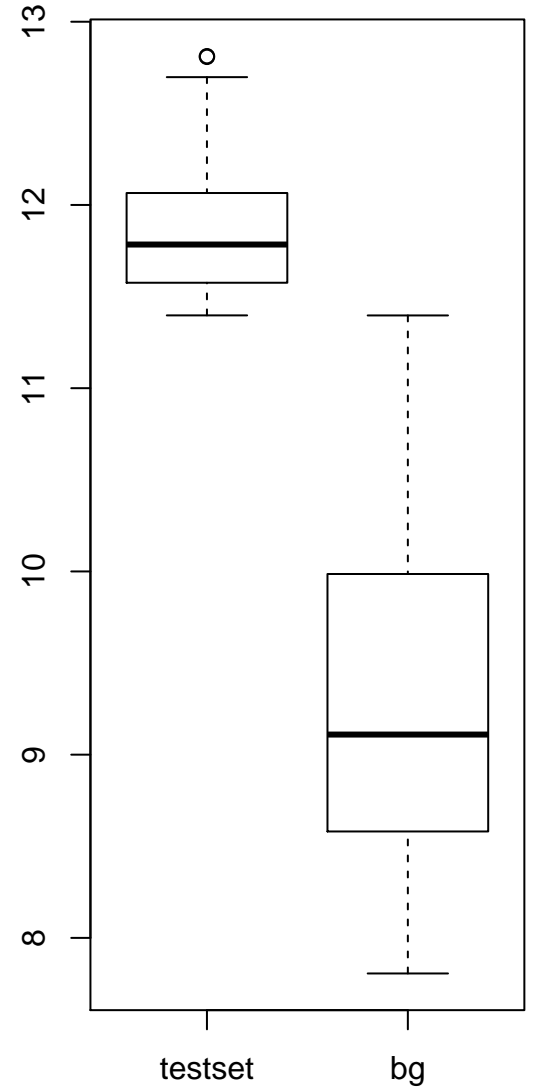
median expression values  
#genes: 1771



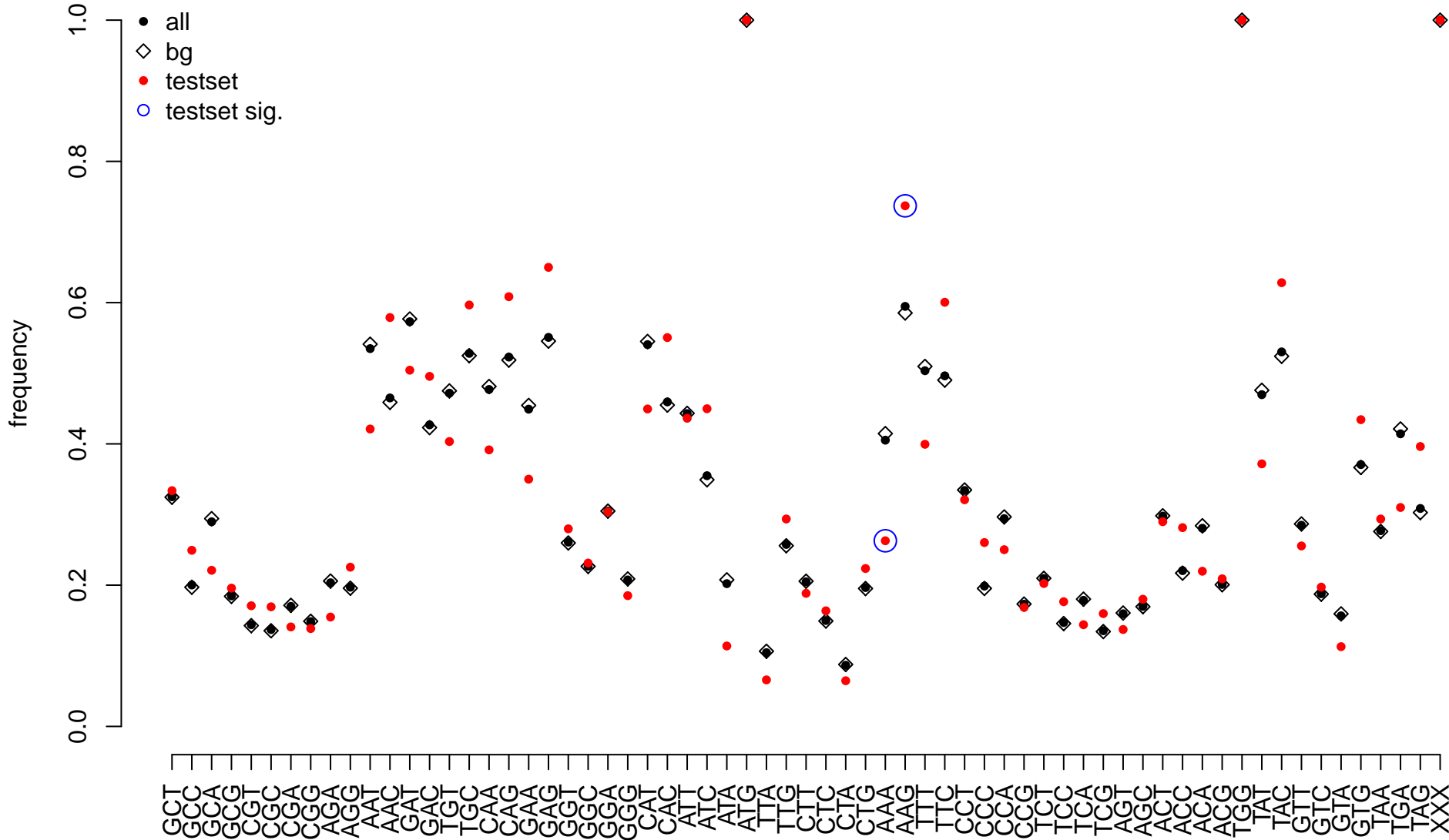
Codon usage frequency spectrum normalized per aa per gene  
group starting from highest expression values #genes: 1869



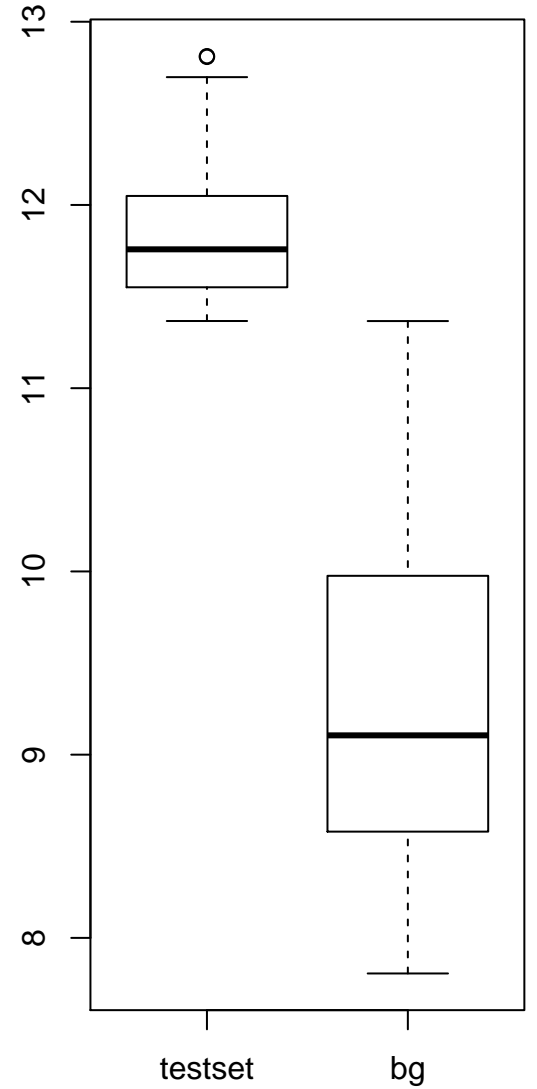
median expression values  
#genes: 1869



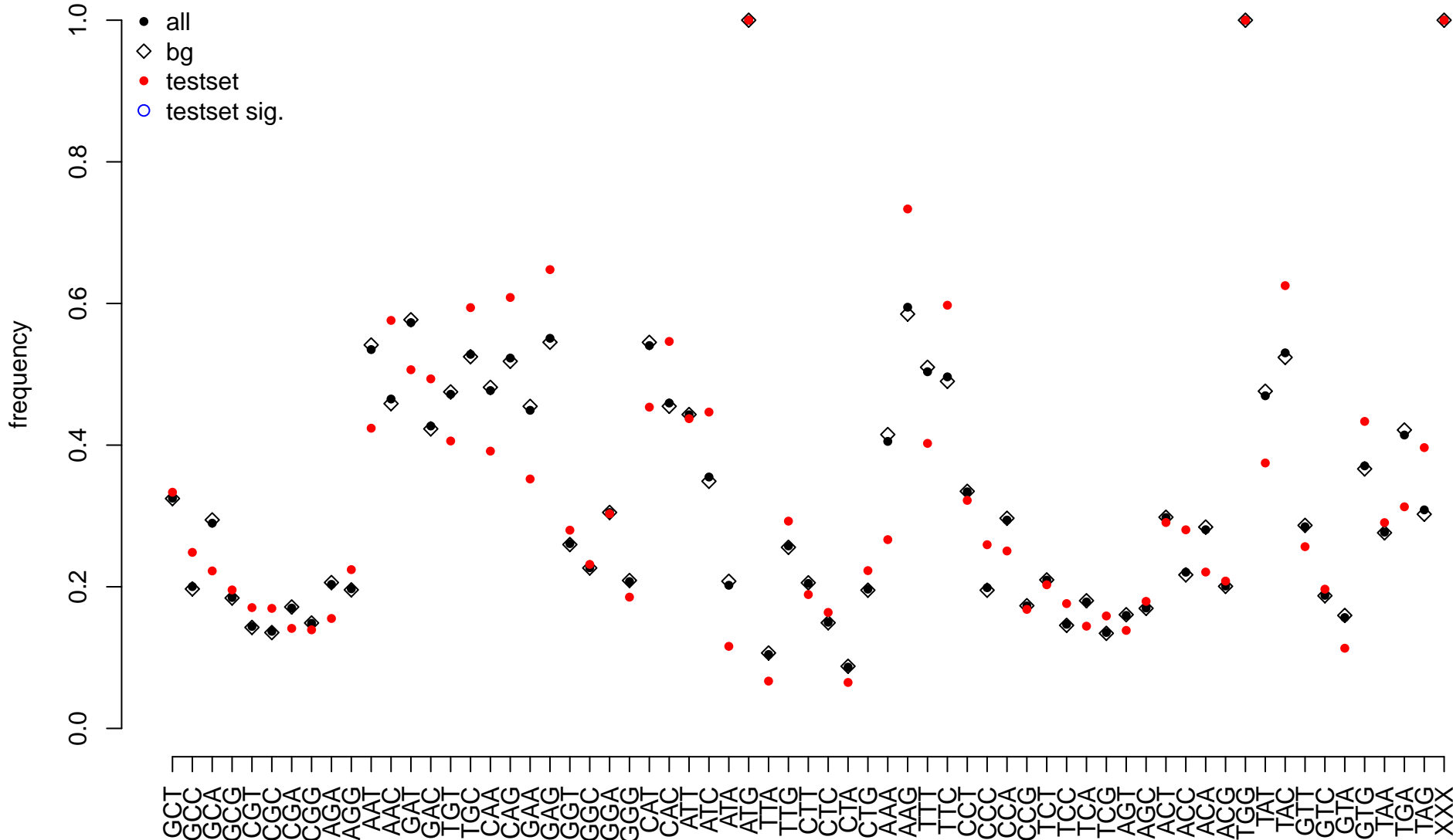
**Codon usage frequency spectrum normalized per aa per gene  
group starting from highest expression values #genes: 1967**



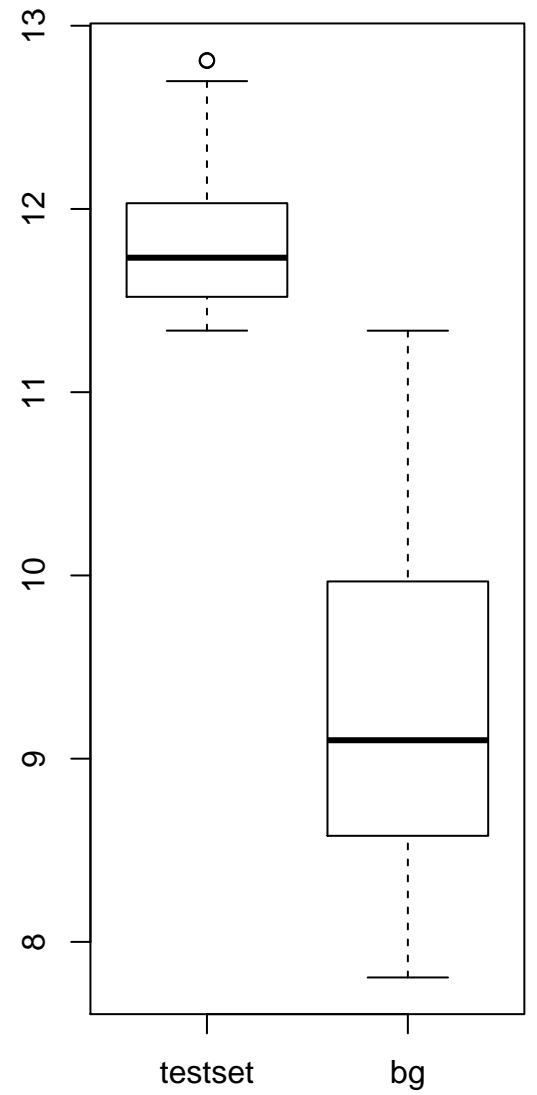
**median expression values  
#genes: 1967**



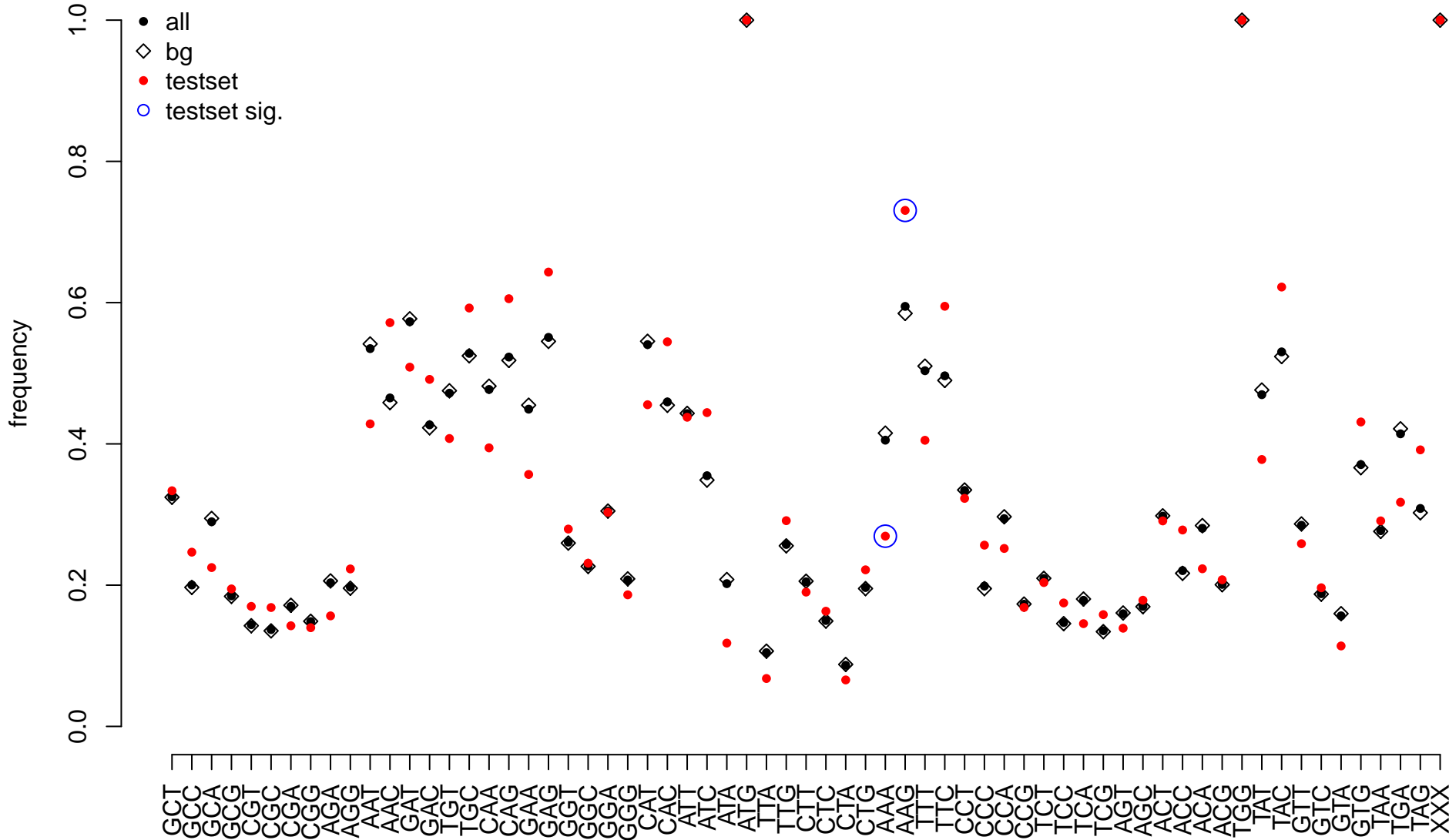
**Codon usage frequency spectrum normalized per aa per gene  
group starting from highest expression values #genes: 2067**



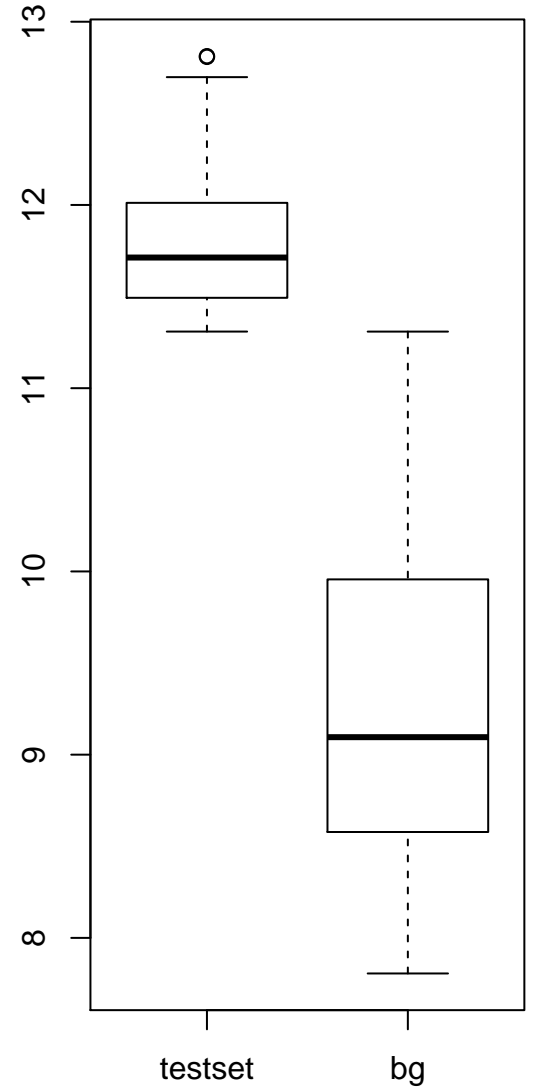
**median expression values  
#genes: 2067**



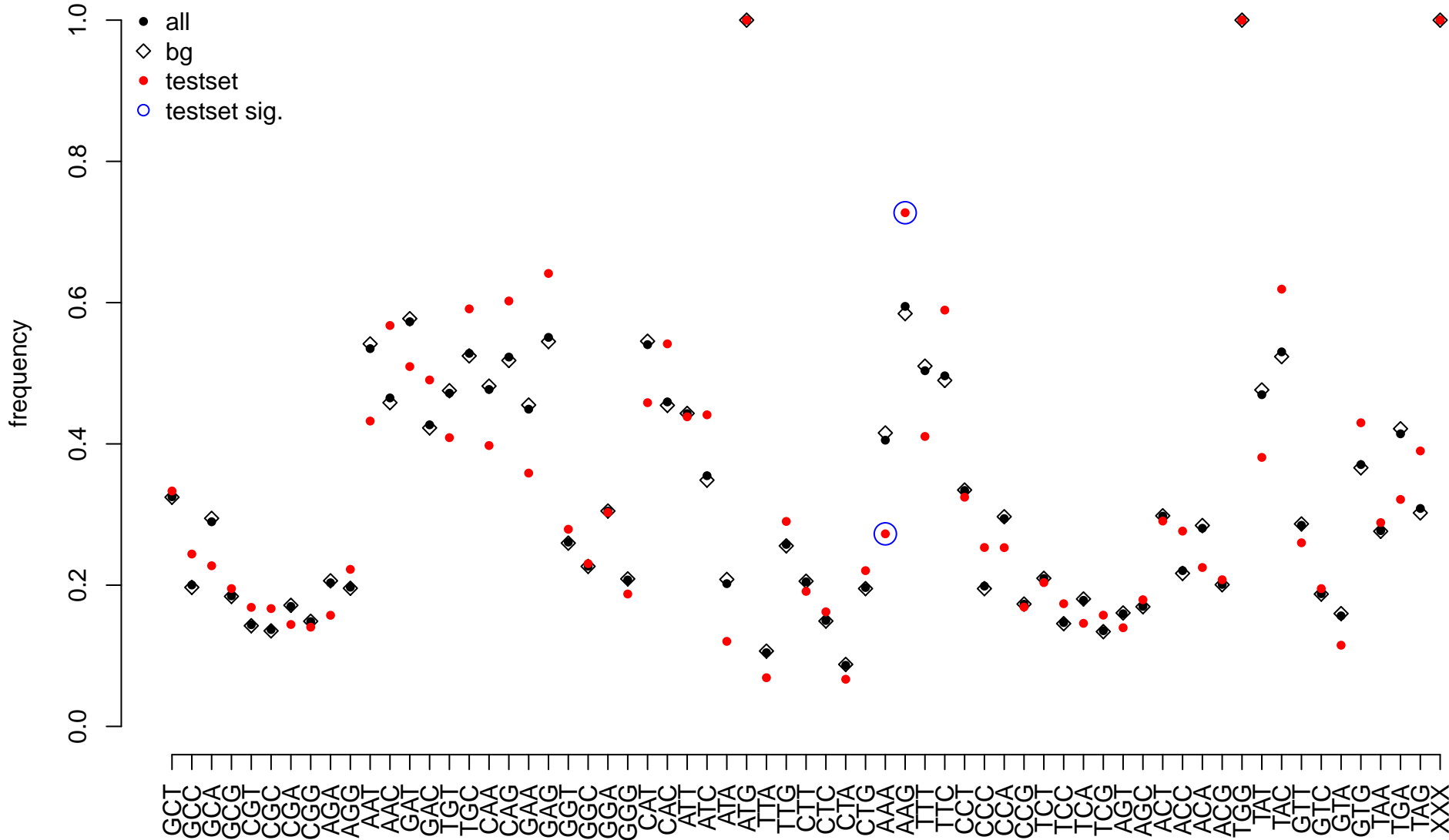
**Codon usage frequency spectrum normalized per aa per gene**  
**group starting from highest expression values #genes: 2167**



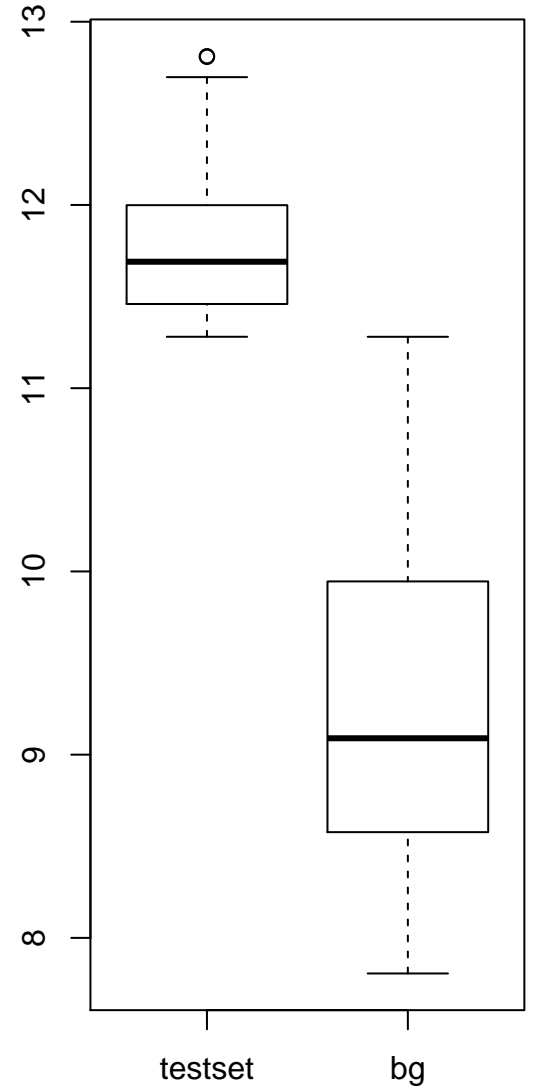
**median expression values**  
**#genes: 2167**



**Codon usage frequency spectrum normalized per aa per gene  
group starting from highest expression values #genes: 2265**

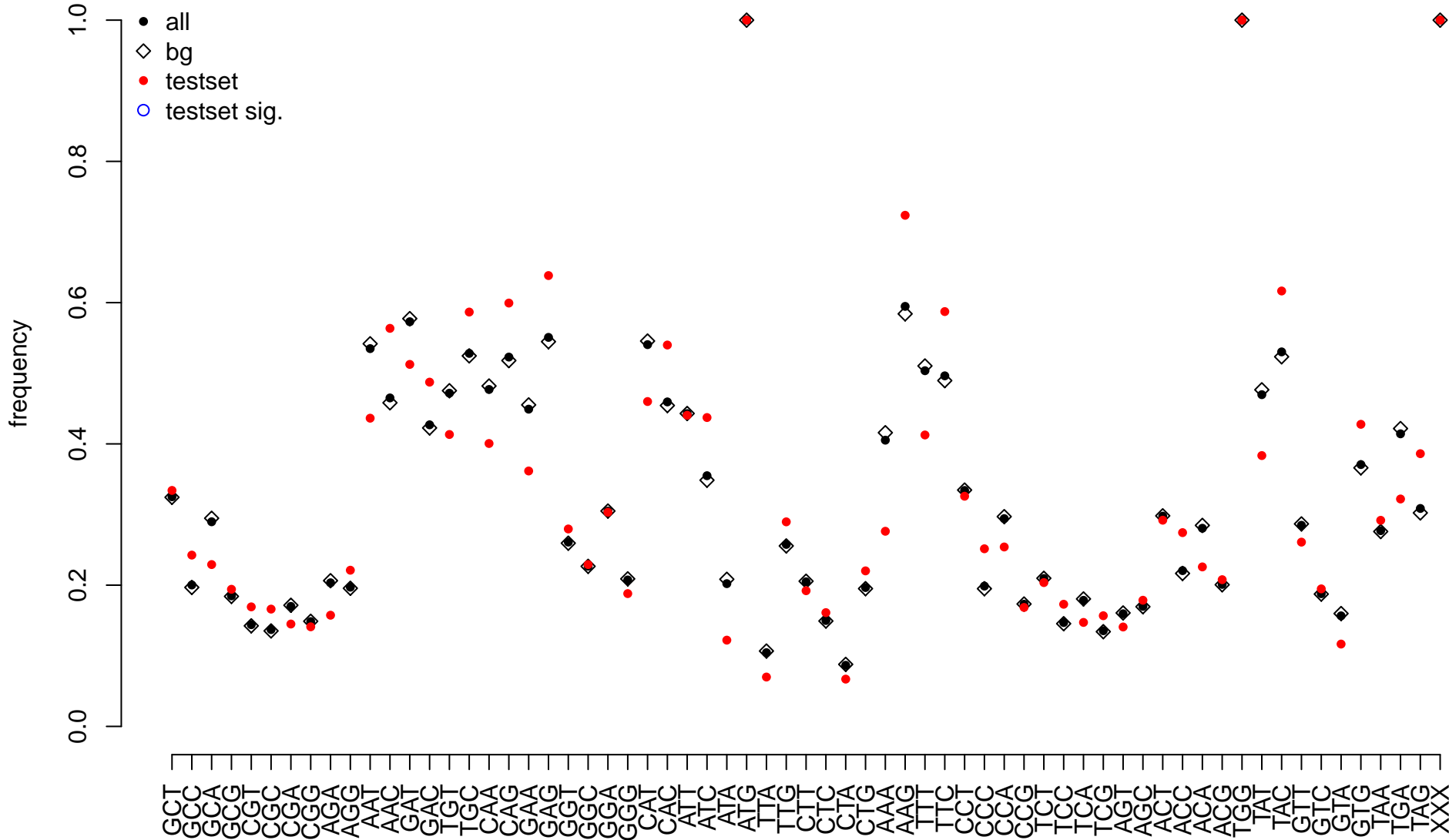


**median expression values  
#genes: 2265**

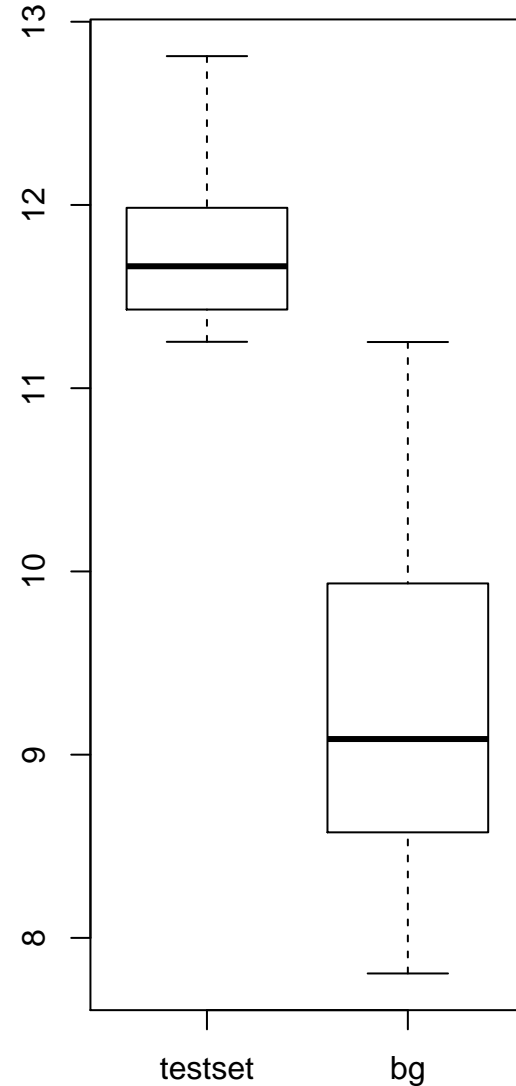




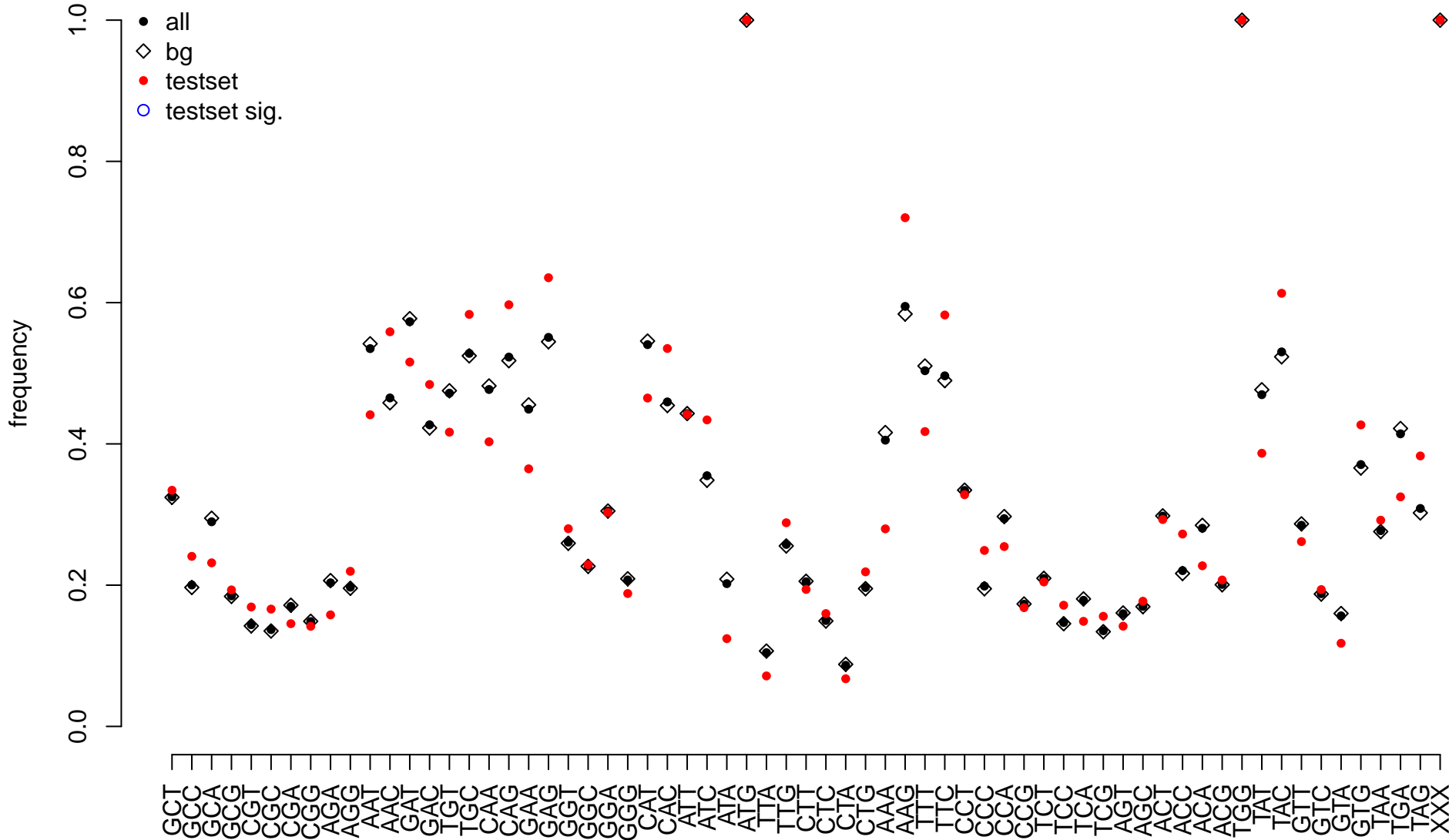
Codon usage frequency spectrum normalized per aa per gene  
group starting from highest expression values #genes: 2363



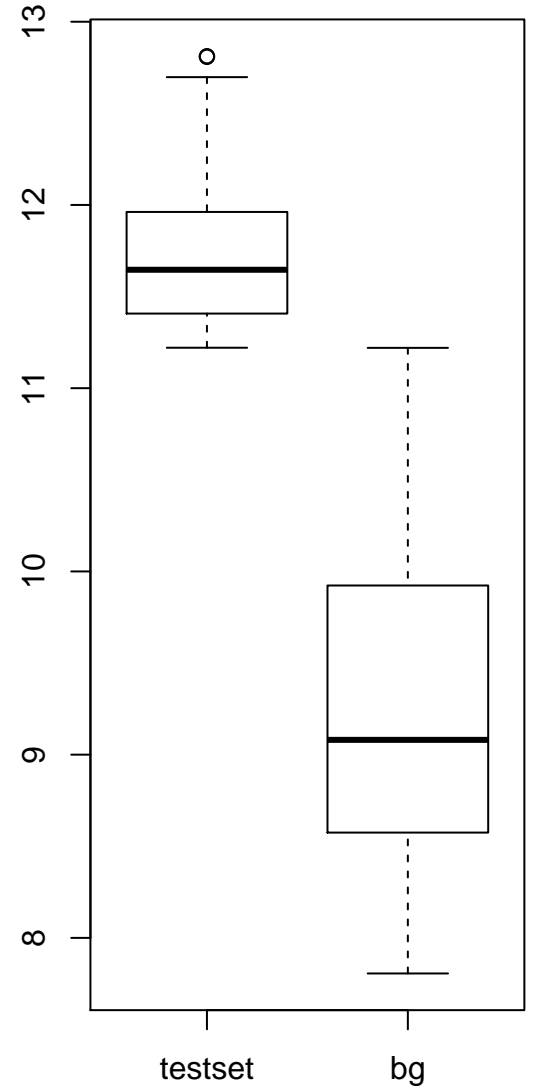
median expression values  
#genes: 2363



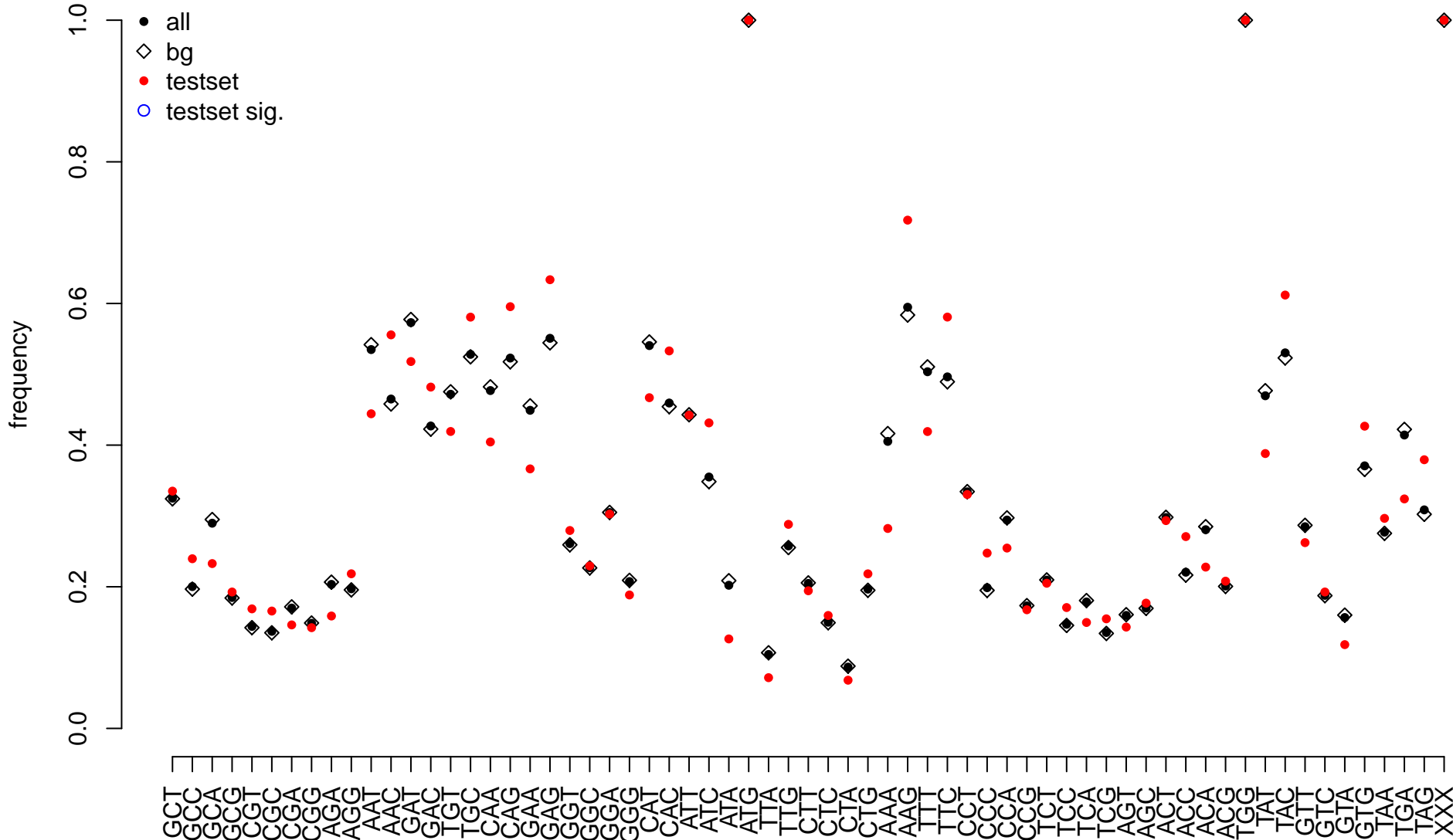
**Codon usage frequency spectrum normalized per aa per gene  
group starting from highest expression values #genes: 2462**



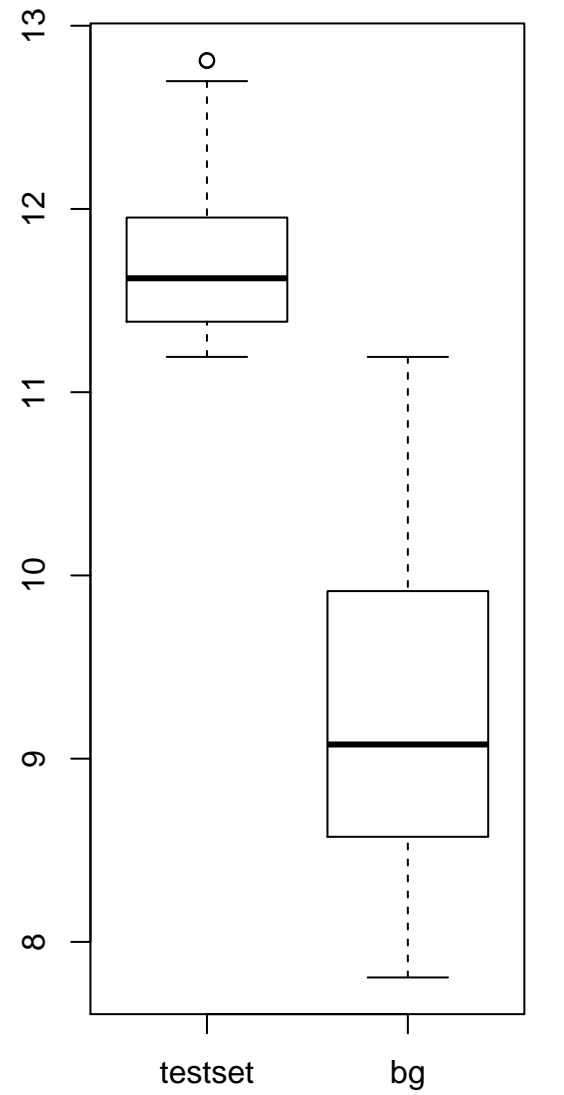
**median expression values  
#genes: 2462**



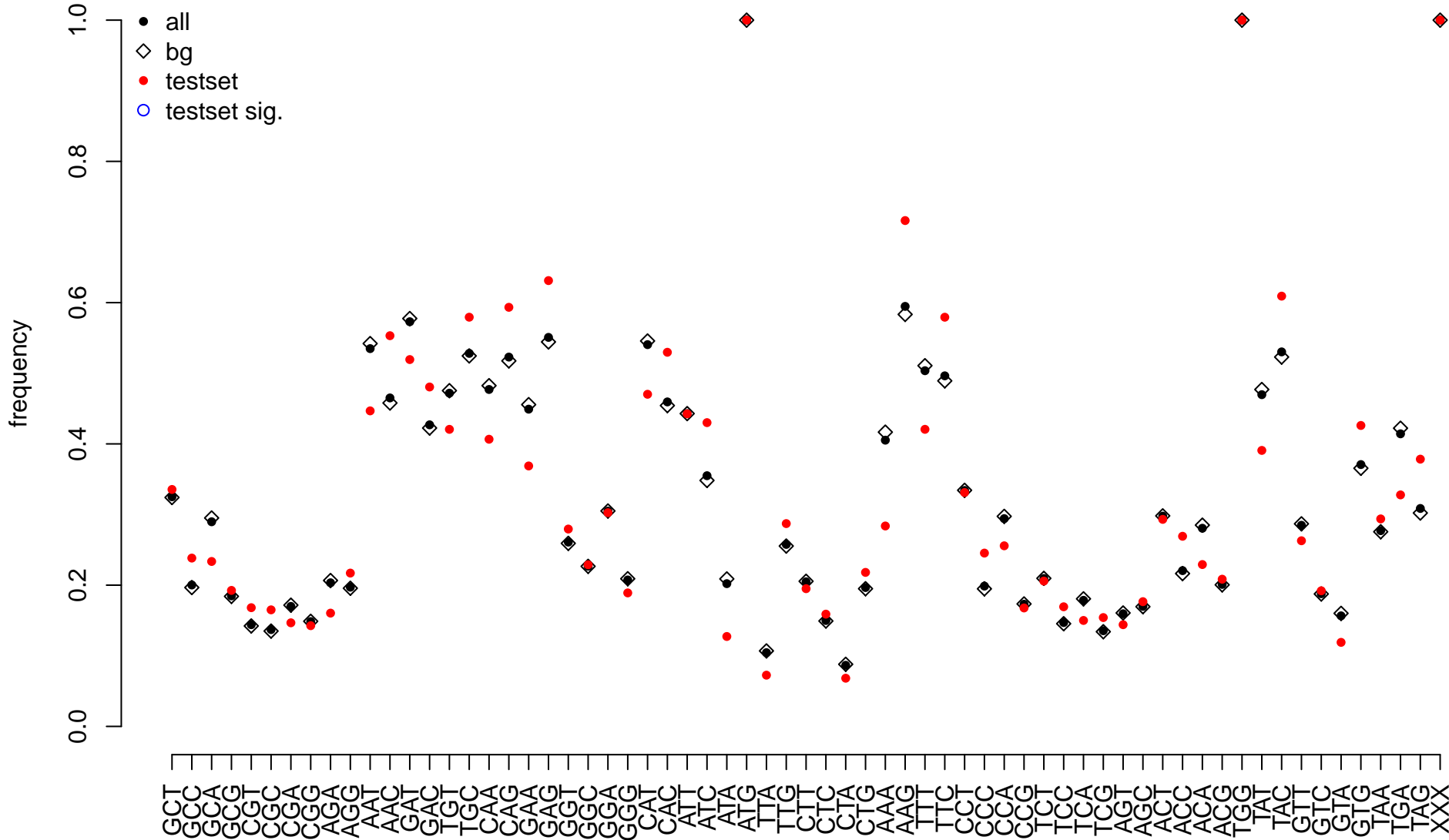
**Codon usage frequency spectrum normalized per aa per gene  
group starting from highest expression values #genes: 2559**



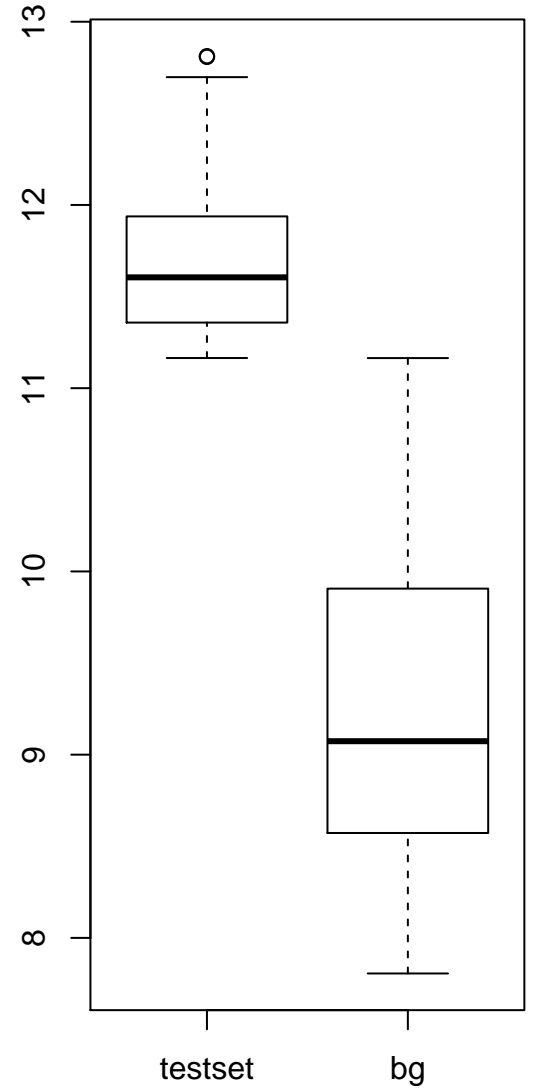
**median expression values  
#genes: 2559**



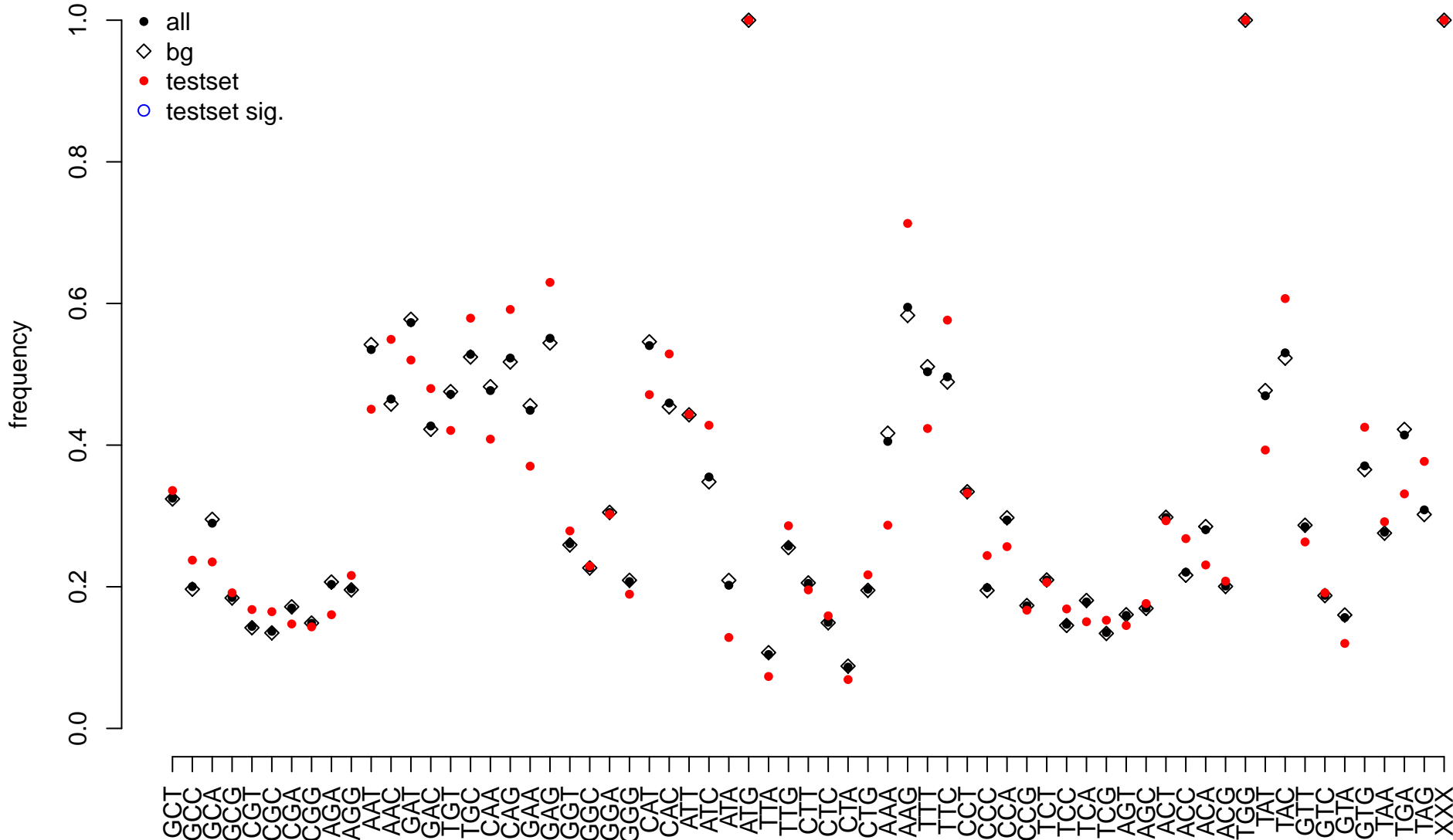
Codon usage frequency spectrum normalized per aa per gene  
group starting from highest expression values #genes: 2658



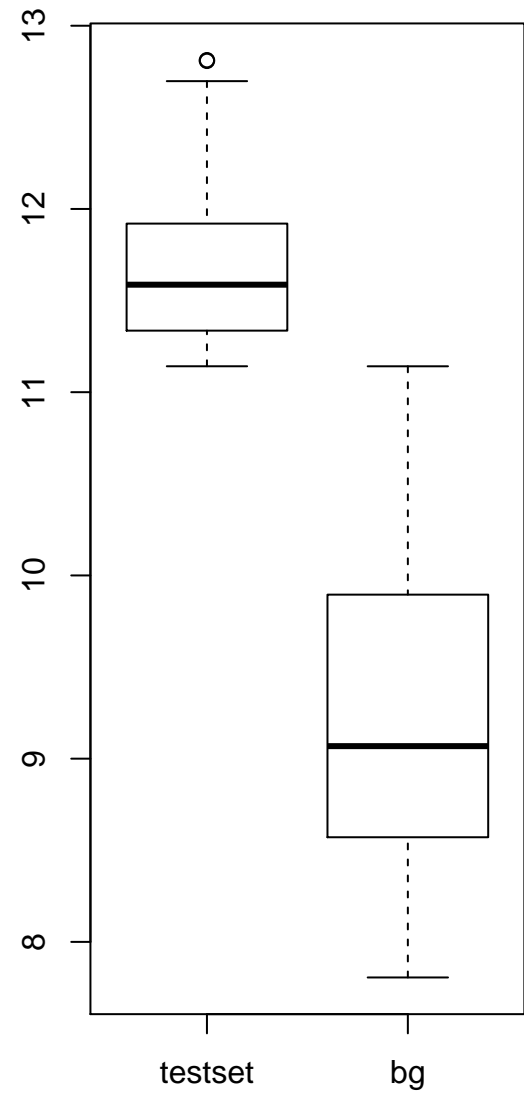
median expression values  
#genes: 2658



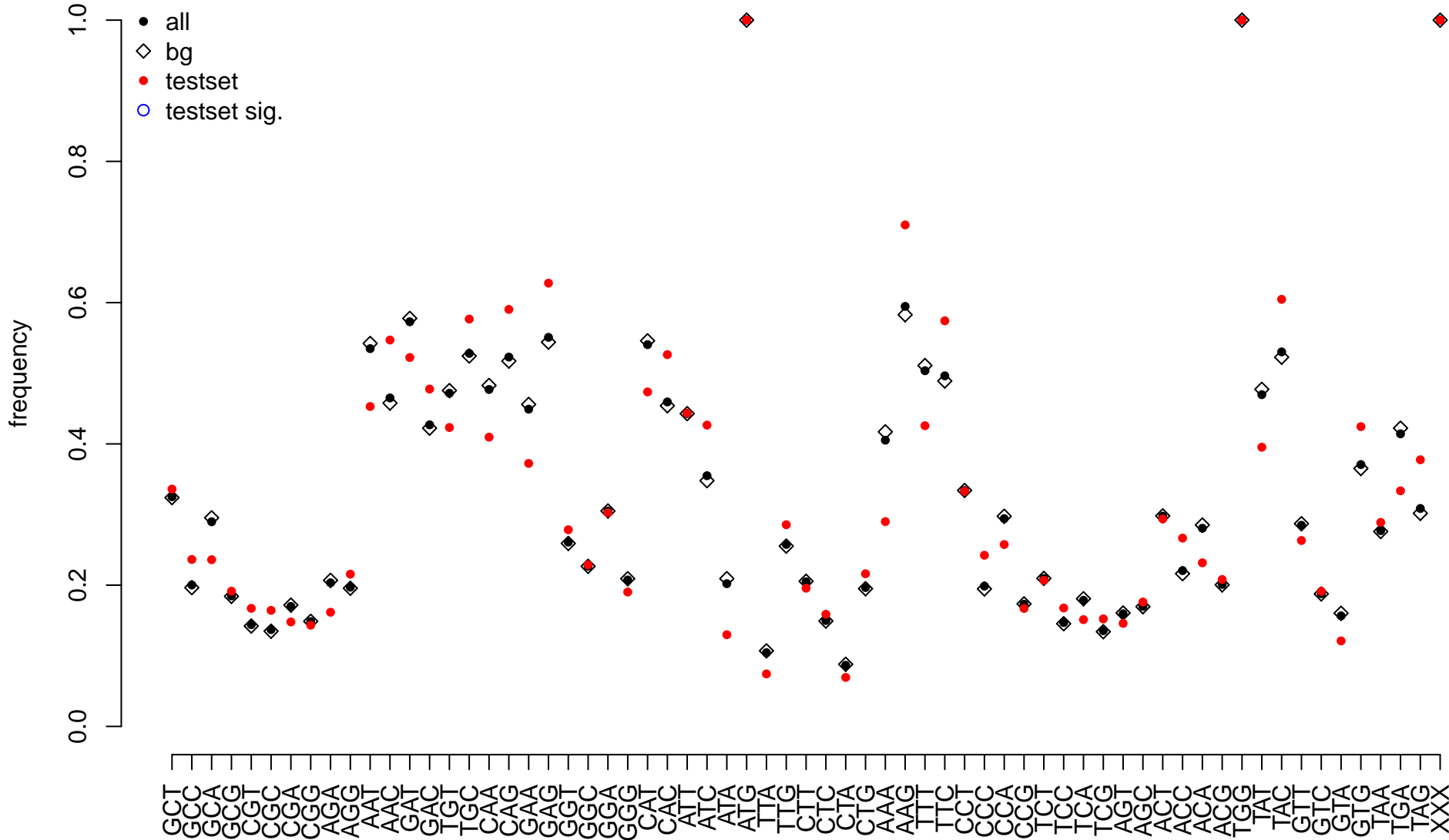
**Codon usage frequency spectrum normalized per aa per gene**  
**group starting from highest expression values #genes: 2757**



**median expression values**  
**#genes: 2757**



**Codon usage frequency spectrum normalized per aa per gene**  
**group starting from highest expression values #genes: 2855**



**median expression values**  
**#genes: 2855**

