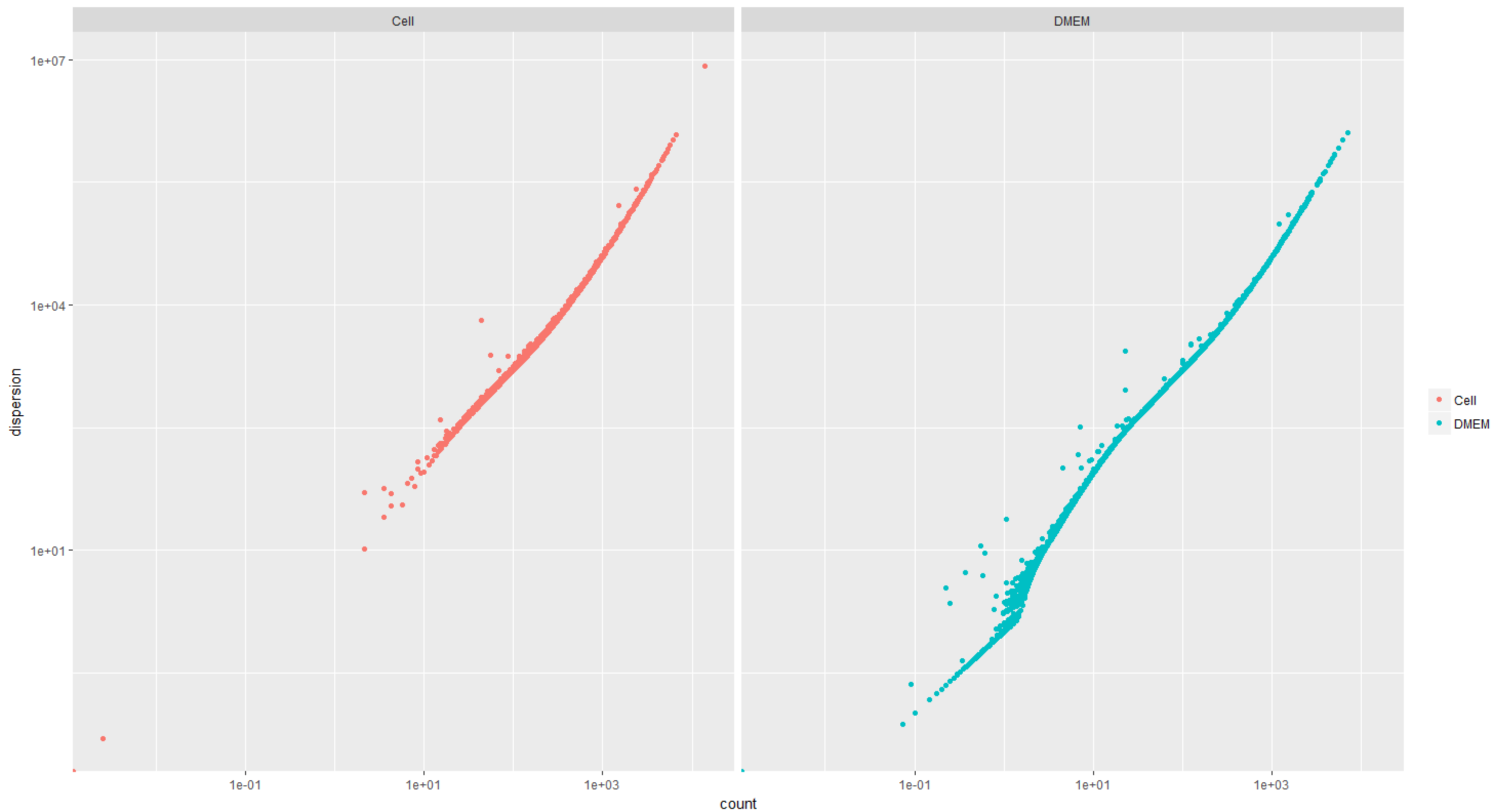
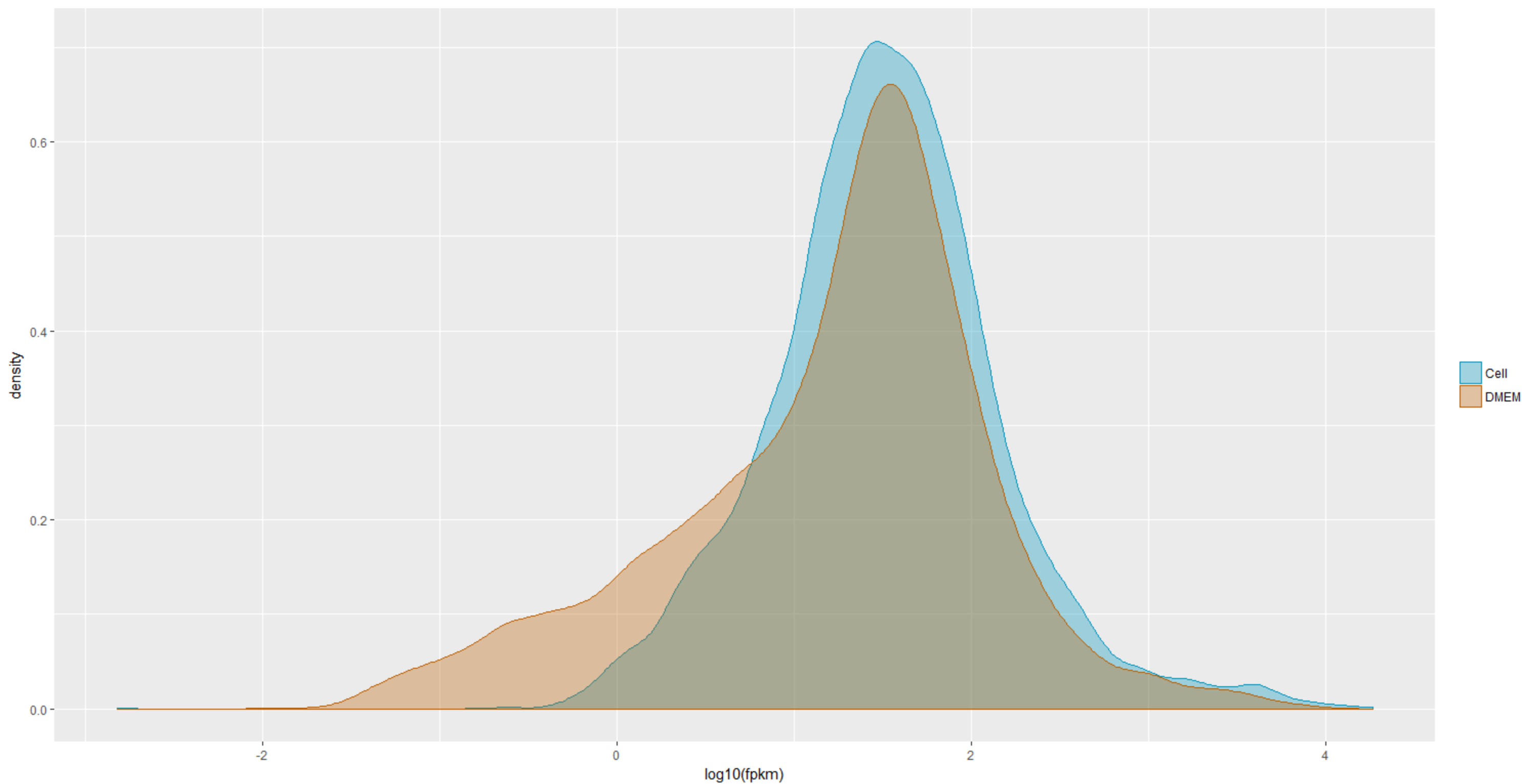


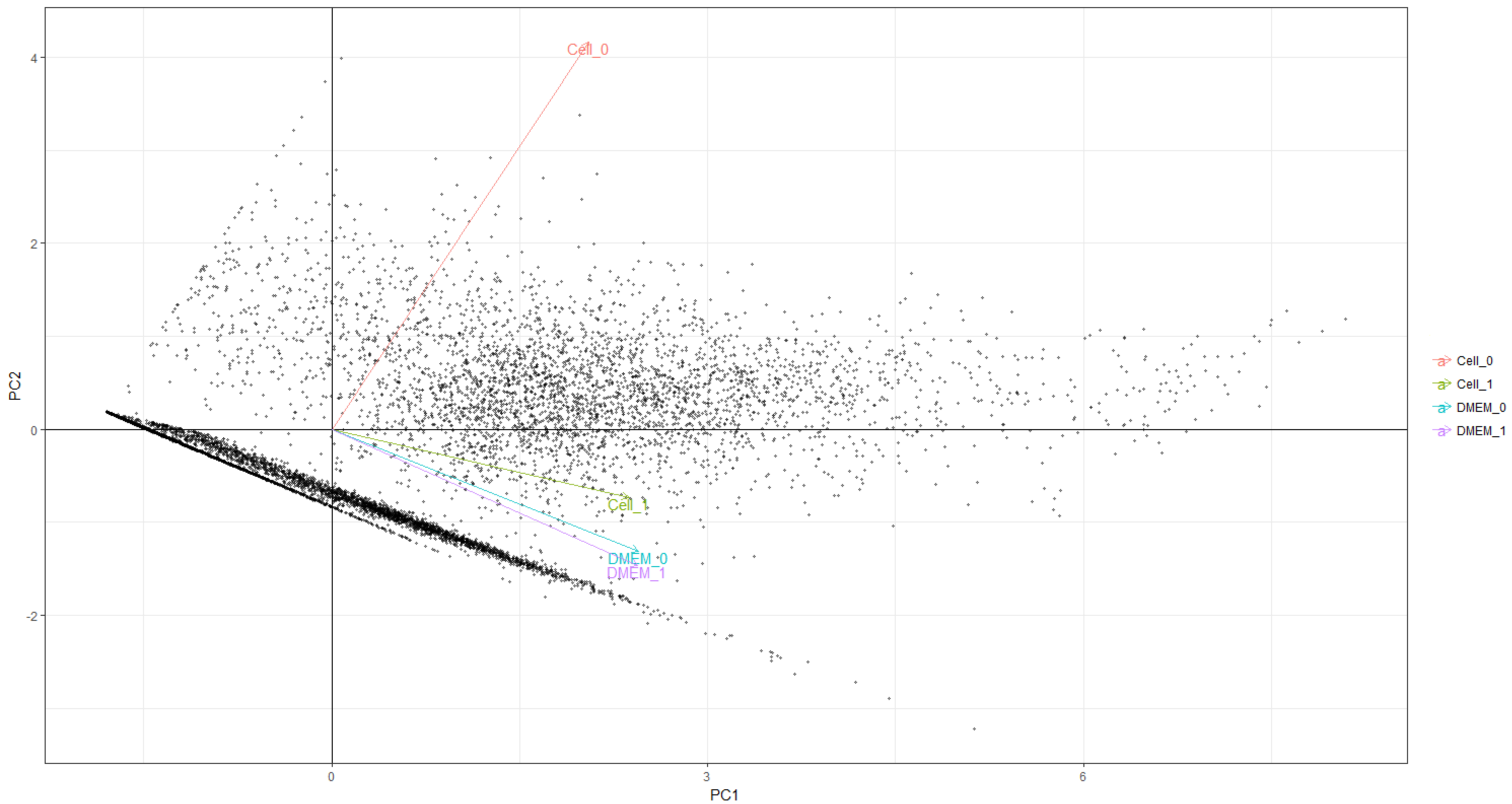


A1: Dispersion graph from *A. fumigatus* in co-culture with A549 cell (Cell) and DMEM

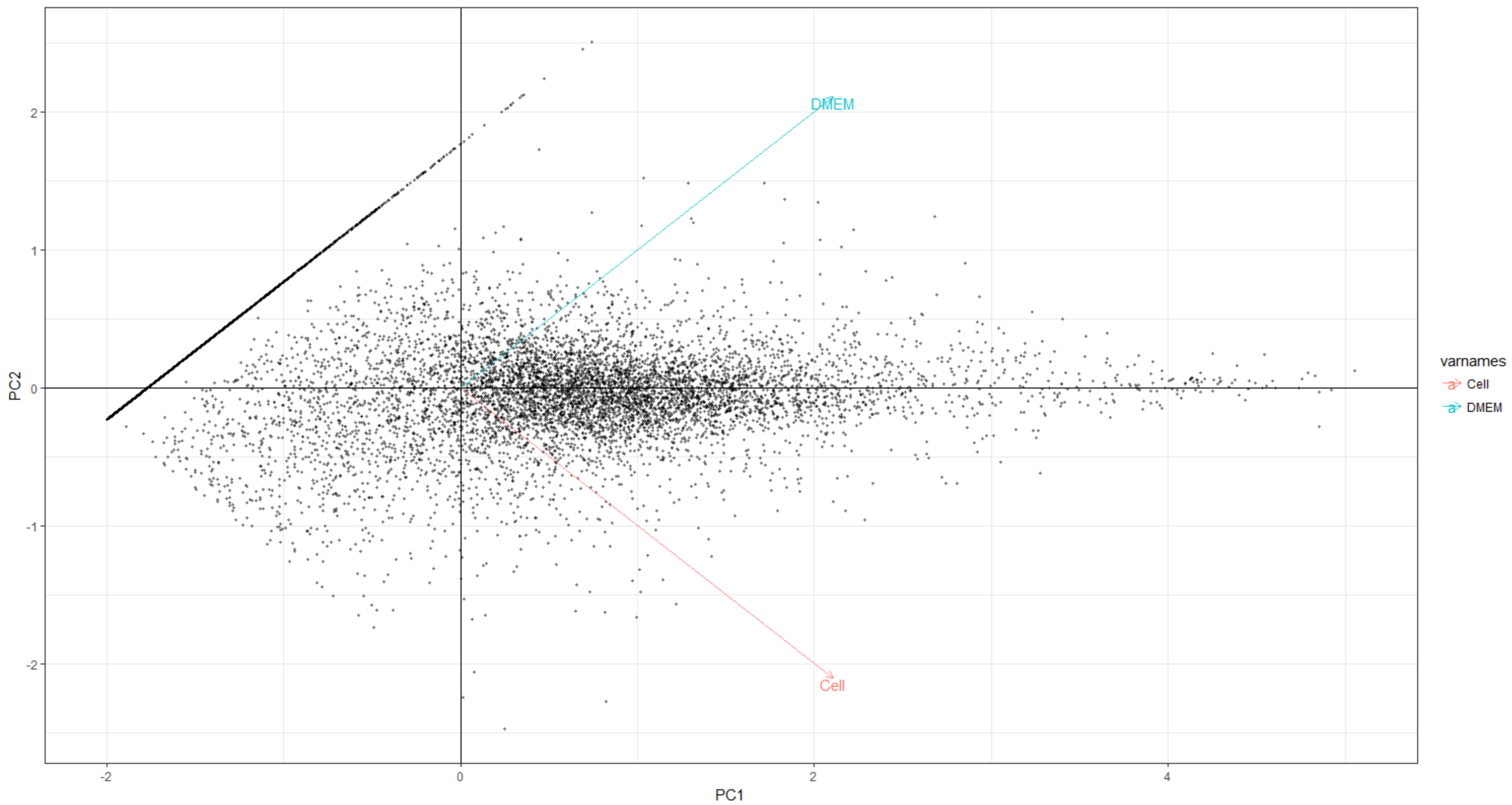


A2: Expression level distribution of mapped genes in *A. fumigatus* in co-culture with A549 cell (Cell) and DMEM. FPKM= fragments per kilobase of transcript per million fragments mapped.





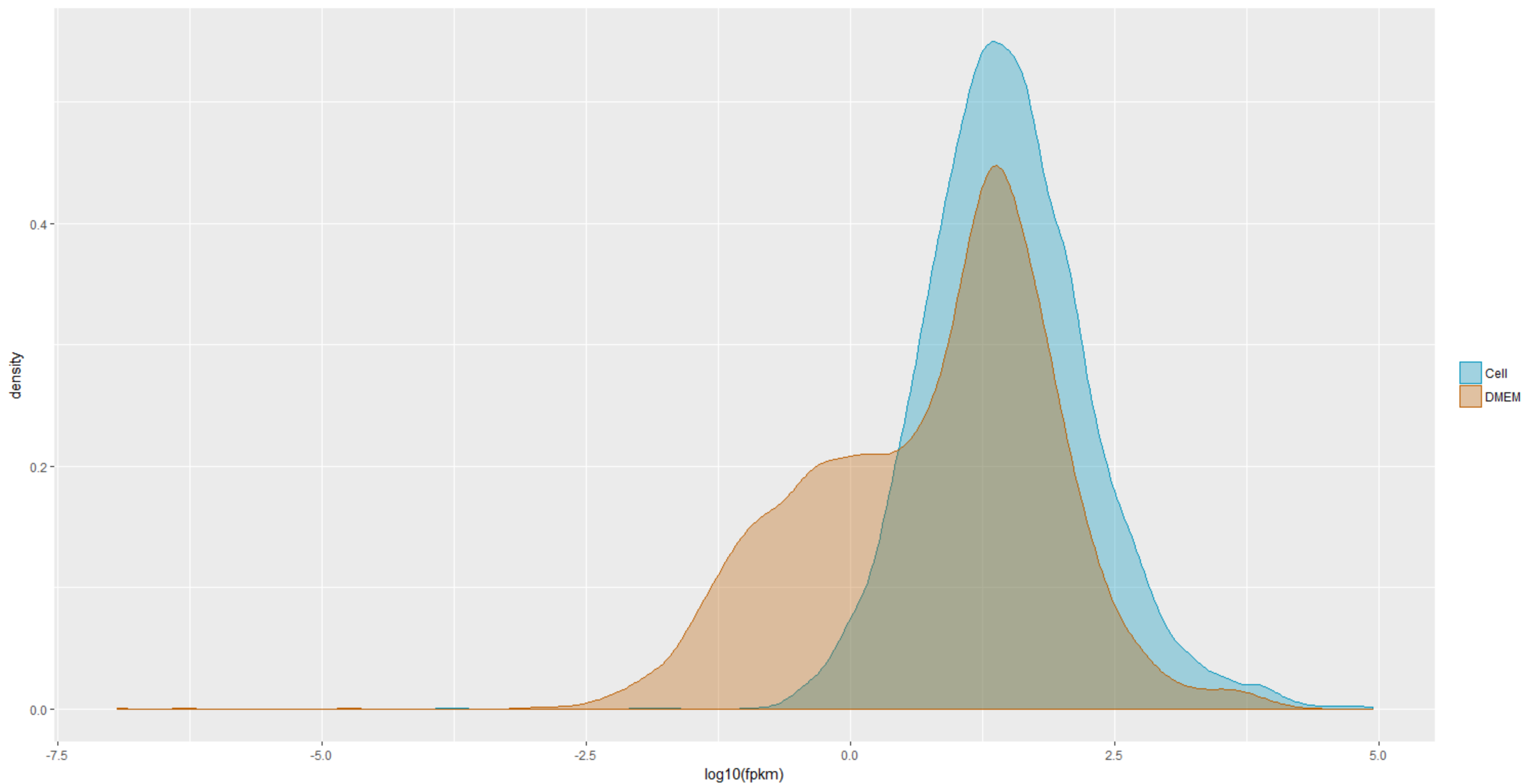
A3: Principal component analysis per sample of RNA-seq data from *A. fumigatus* in co-culture with A549 cell (Cell) and DMEM.



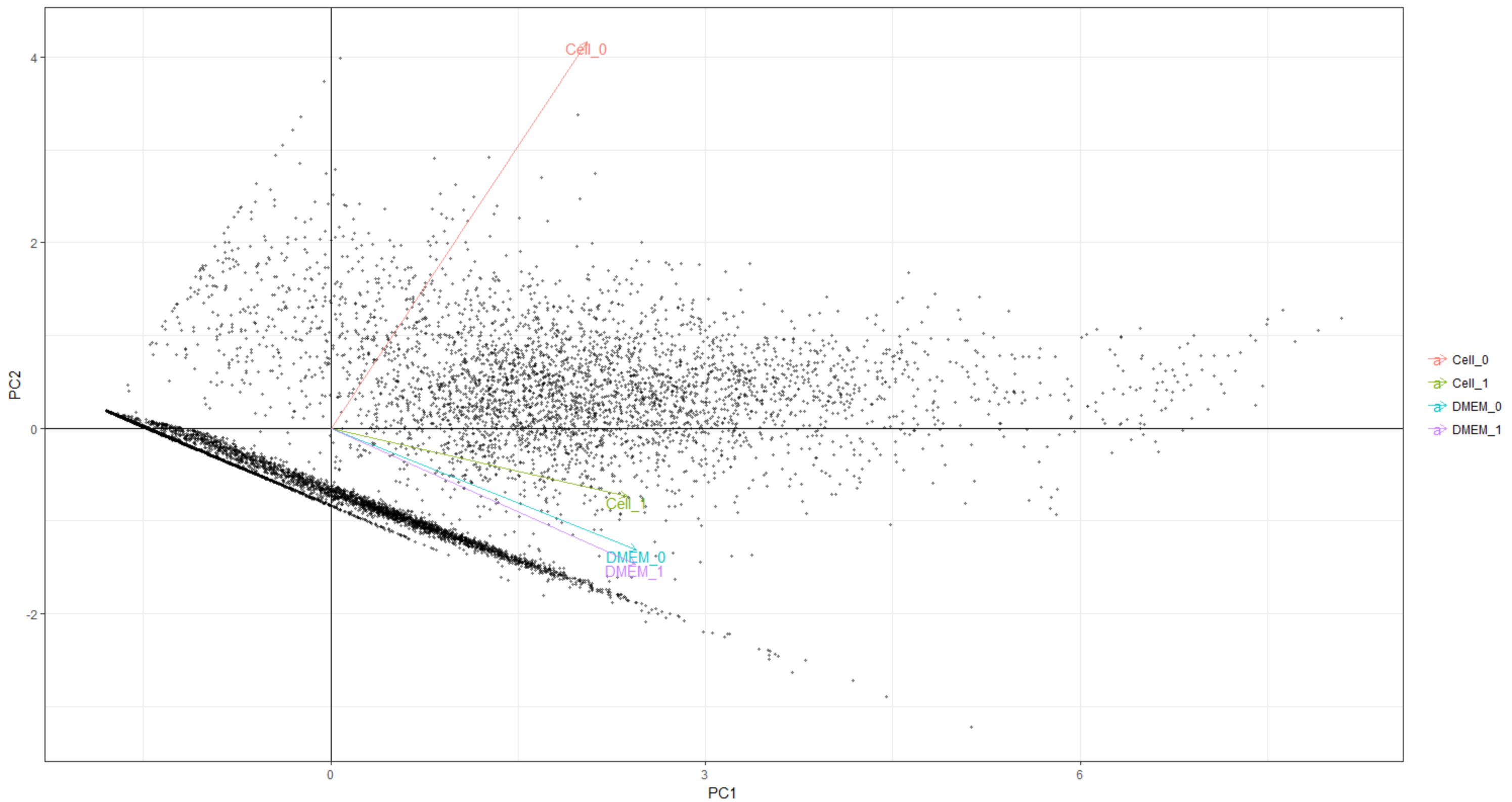
A4: Principal component analysis of RNA-seq data from *A. fumigatus* in co-culture with A549 cell (Cell) and DMEM



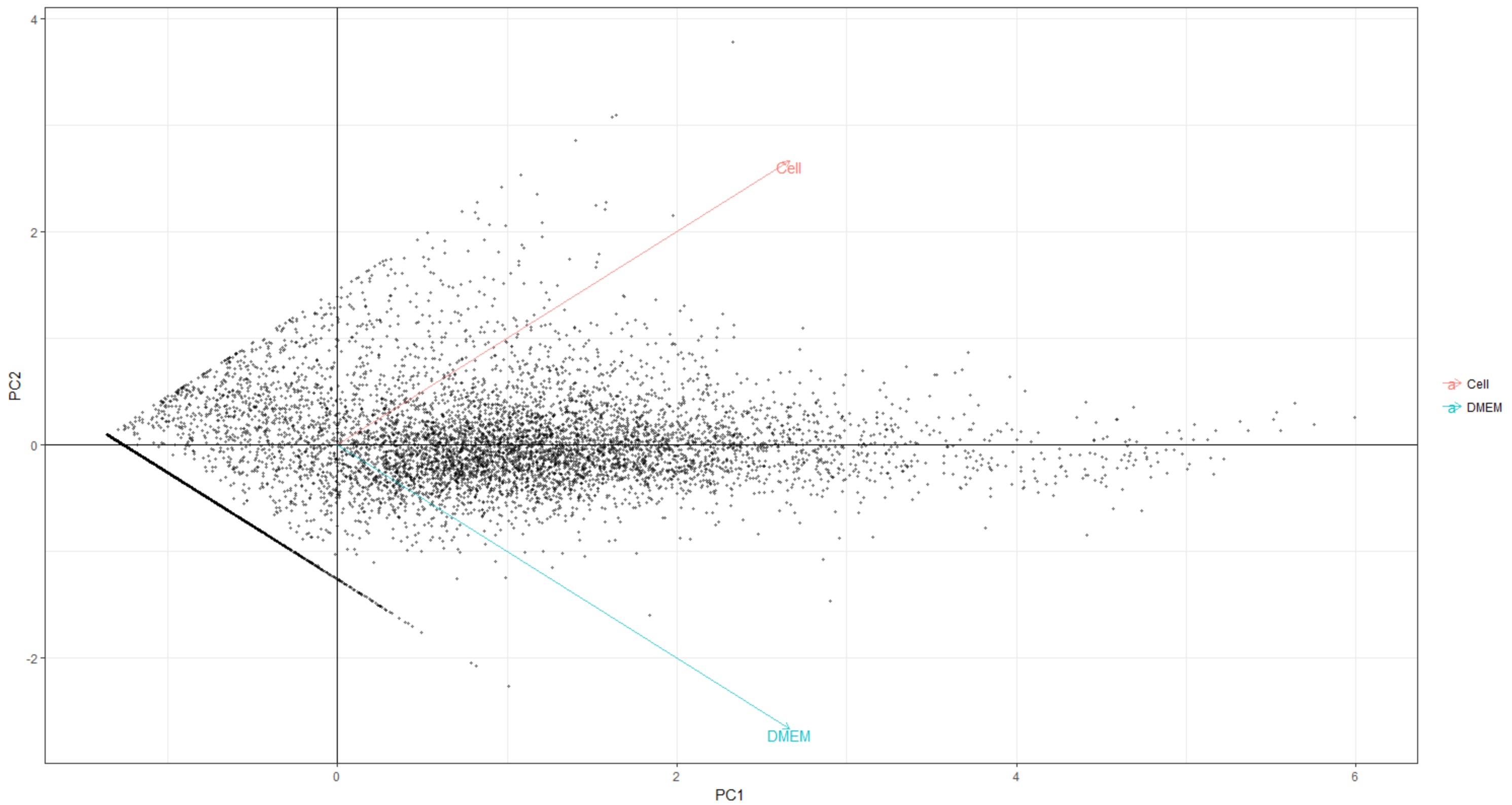
B1: Dispersion graph from *A. niger* in co-culture with A549 cell (Cell) and DMEM.



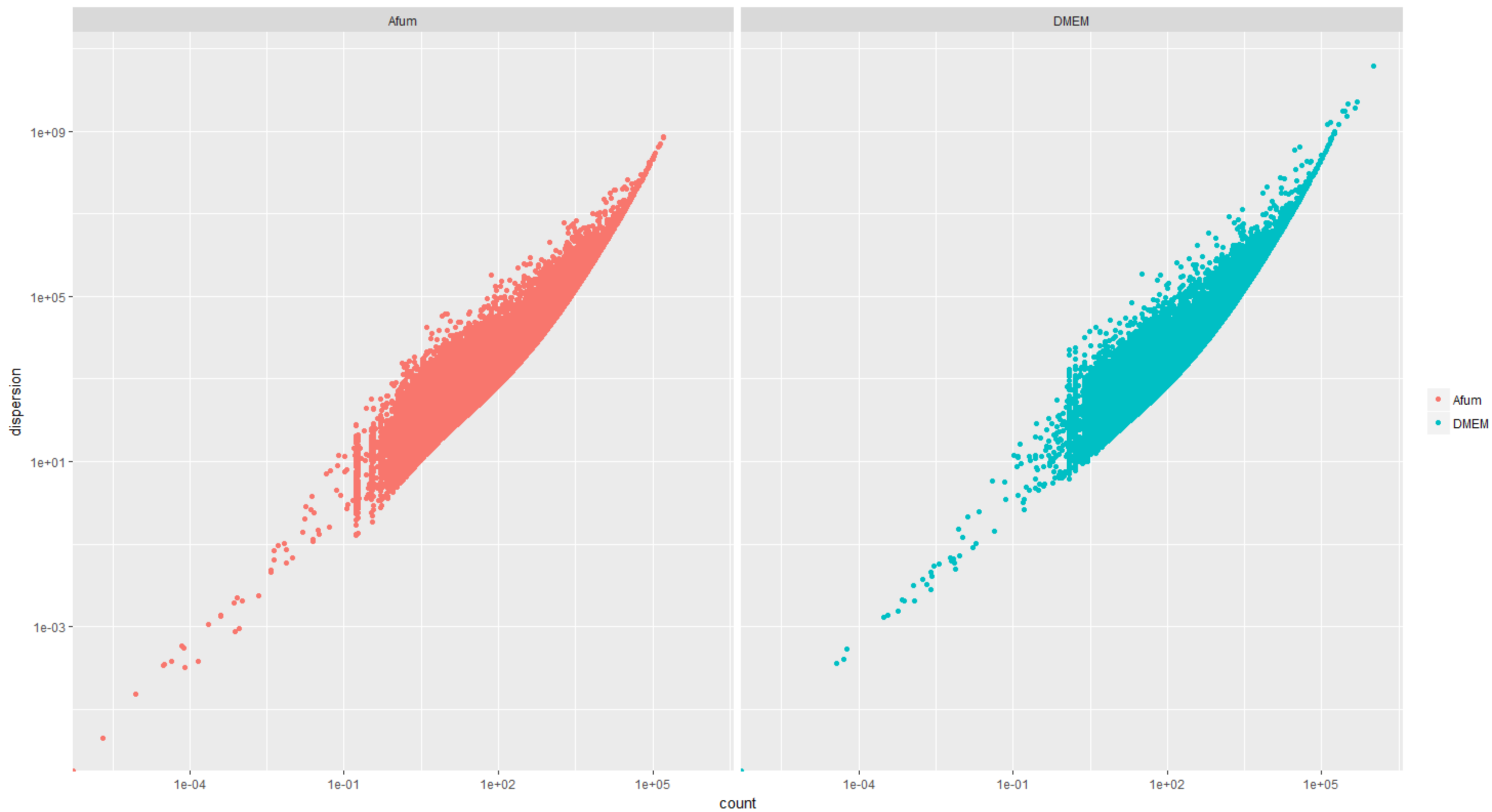
B2: Expression level distribution of mapped genes in *A. niger* in co-culture with A549 cell (Cell) and DMEM. FPKM= fragments per kilobase of transcript per million fragments mapped.



B3: Principal component analysis per sample of RNA-seq data from *A. niger* in co-culture with A549 cell (Cell) and DMEM.

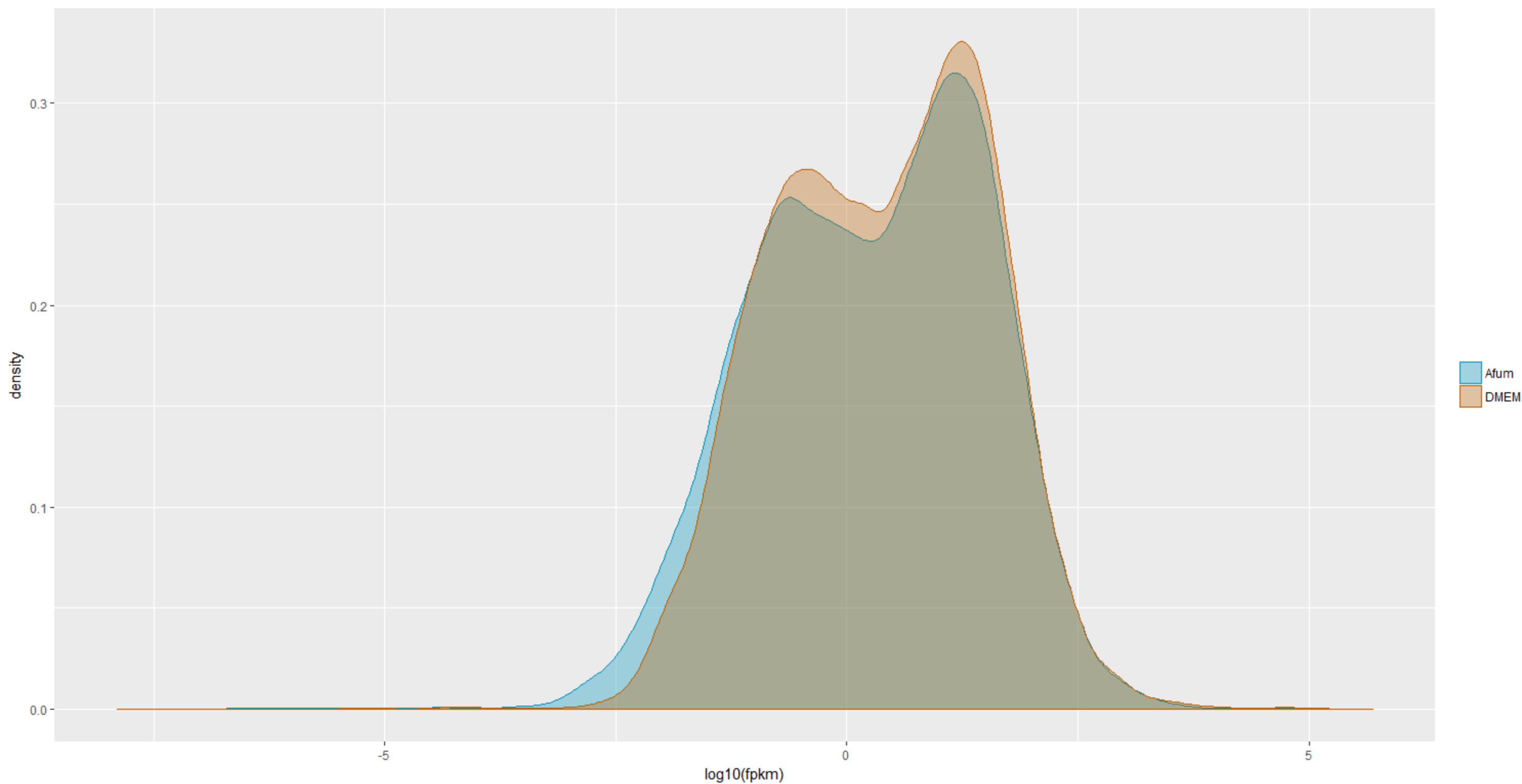


B4: Principal component analysis of RNA-seq data from *A. niger* in co-culture with A549 cell (Cell) and DMEM.



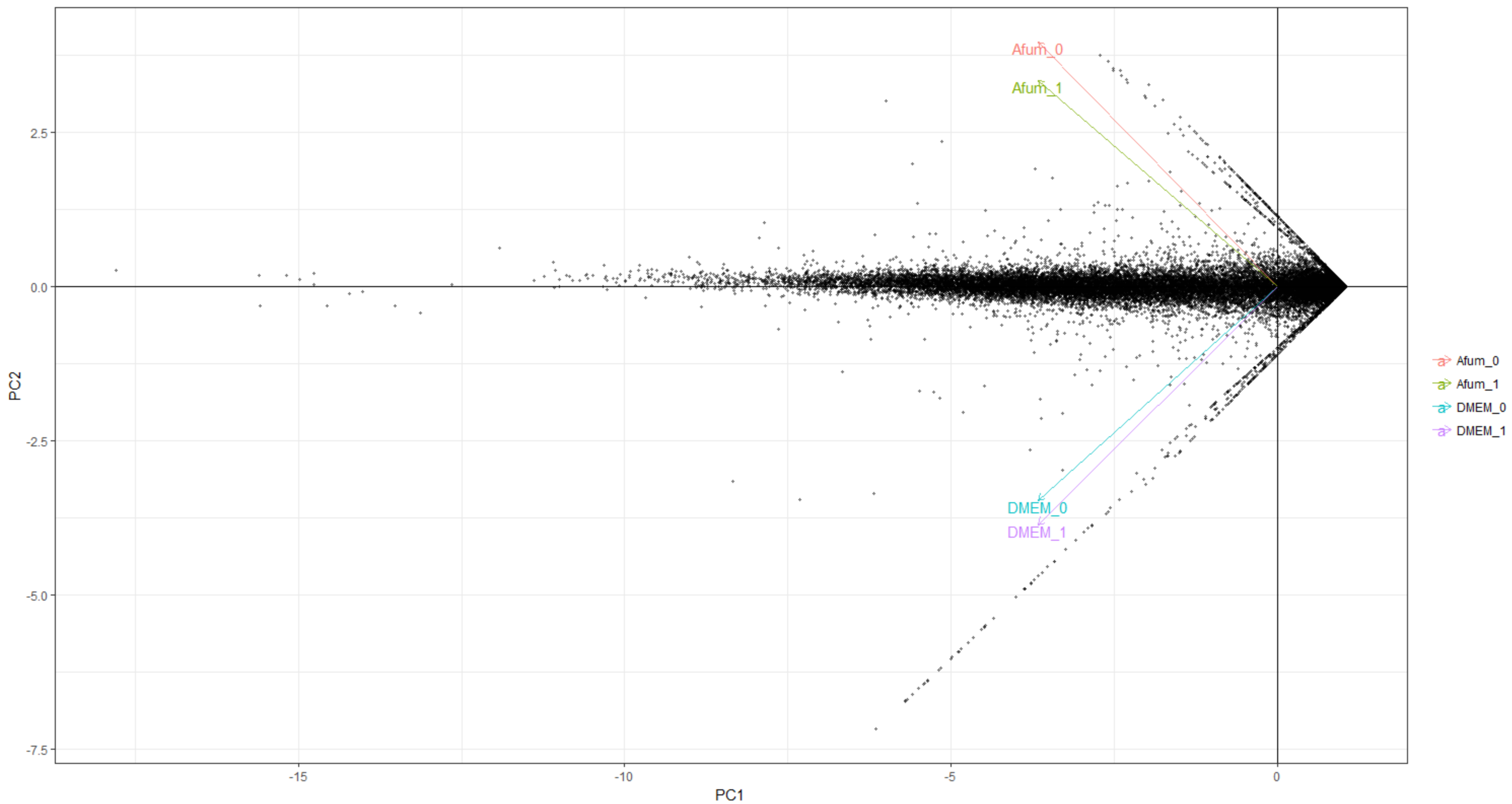
C1: Dispersion graph from A549 cells in co-culture with *A. fumigatus* (Afum) and DMEM.



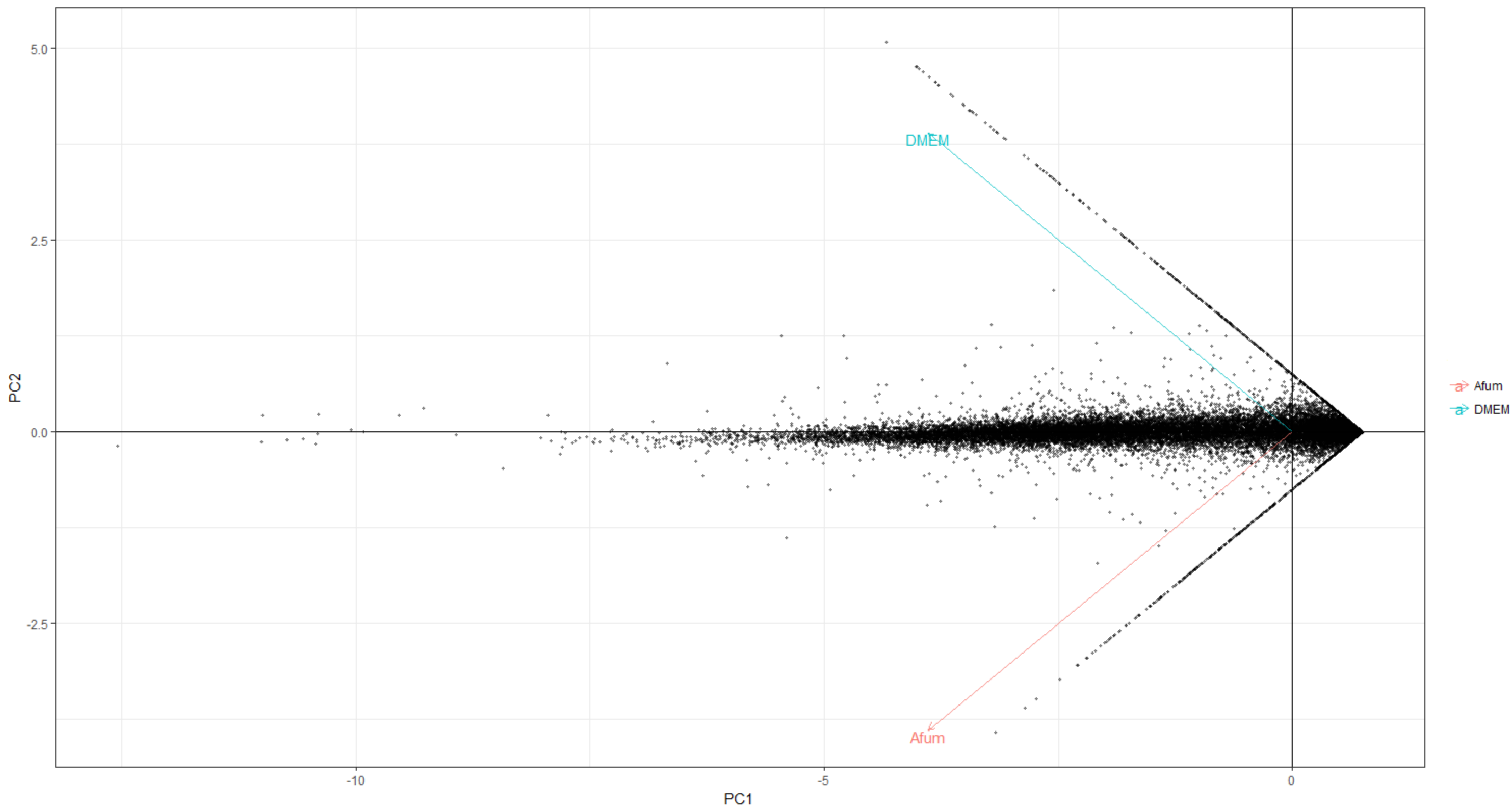


C2: Expression level distribution of mapped genes in A549 cells in co-culture with *A. fumigatus* (Afum) and DMEM. FPKM= fragments per kilobase of transcript per million fragments mapped.





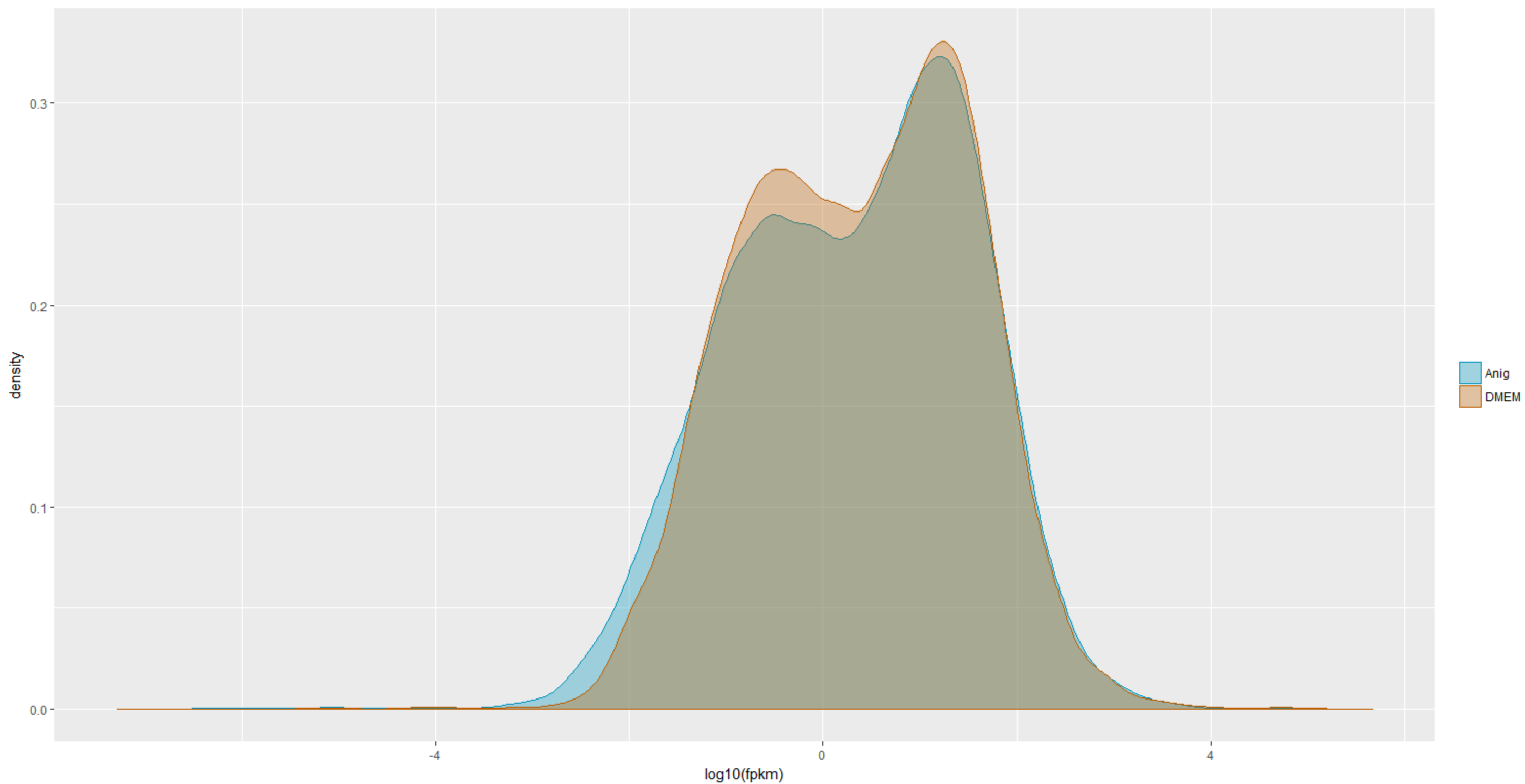
C3: Principal component analysis per sample of RNA-seq data from A549 cells in co-culture with *A. fumigatus* (Afum) and DMEM.



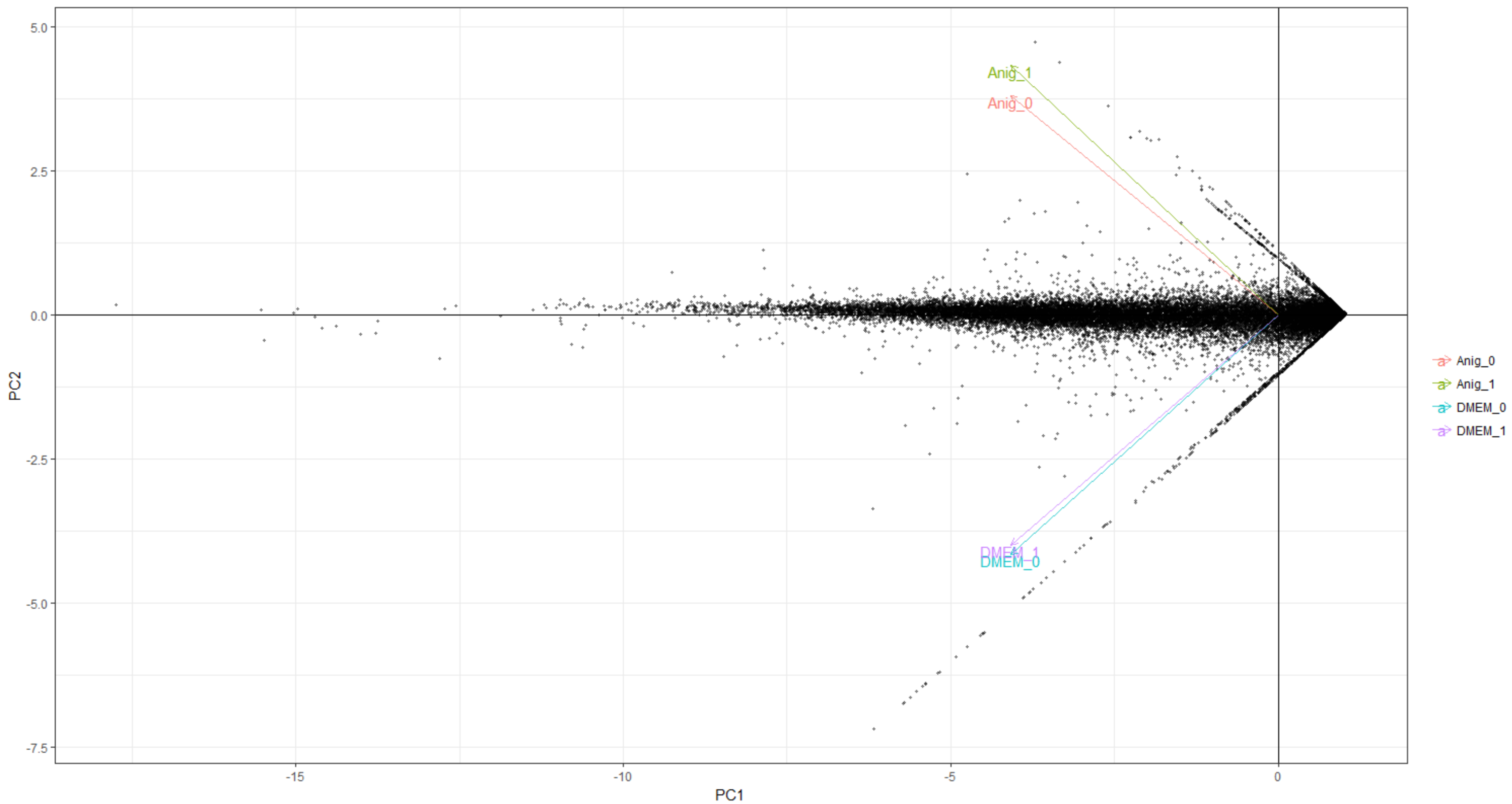
C4: Principal component analysis of RNA-seq data from A549 cells in co-culture with *A. fumigatus* (Afum) and DMEM.



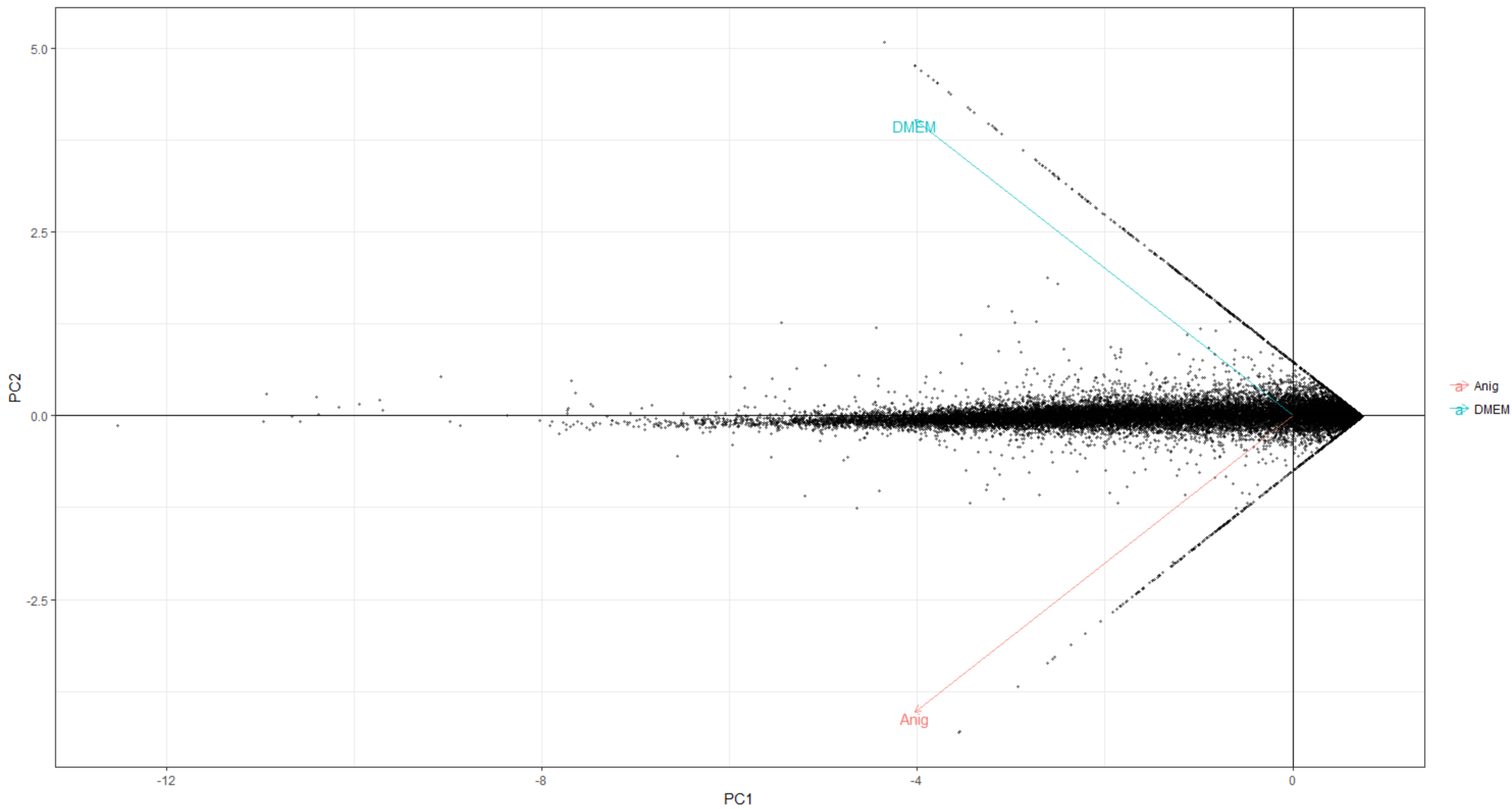
D1: Dispersion graph from A549 cells in co-culture with *A. niger* (Anig) and DMEM.



D2: Expression level distribution of mapped genes in A549 cells in co-culture with *A. niger* (Anig) and DMEM. FPKM= fragments per kilobase of transcript per million fragments mapped.



D3: Principal component analysis per sample of RNA-seq data from A549 cells in co-culture with *A. niger* (Anig) and DMEM.



D4: Principal component analysis of RNA-seq data from A549 cells in co-culture with *A. niger* (Anig) and DMEM.