```
In [41]: dfgroupbymean_total_corr = dfgroupby.mean().loc[idx[:,:,:,:],'total_corr'].unstack(level='algorithm_name')
In [42]: dfgroupbymean total corr[ dfgroupbymean total corr.idxmin(axis='columns') == 'QICA']
algorithm name
                                          GLICA
                                                    OICA
                                                           america
                                                                      sa4ica
prime dimension n samples distribution
                         pure zipf
                                       0.022744 0.016832 0.022744 0.022744
               8192
                16384
                         pure zipf
                                       0.023388 0.017228 0.023388 0.023388
```