['climate', 'P300\_25', 'P1000\_30', 'PMG'] **(4, 34)**  
['climate', 'P600\_25', 'P1000\_30', 'lipids'] **(4, 32)**  
['genotype', 'P600\_25', 'PMG'] **(3, 34)**  
['genotype', 'P300\_25', 'P600\_25', 'P1000\_30', 'PMG'] **(5, 33)**  
['genotype', 'P300\_25', 'P600\_25', 'P1000\_30', 'lipids'] **(5, 29)**  
['P300\_25', 'P600\_25', 'P1000\_30', 'seed\_yield'] **(4, 24)**  
['P300\_25', 'P1000\_30', 'harvest\_25', 'PMG'] **(4, 24)**  
['P300\_25', 'P600\_25', 'P1000\_30', 'proteins'] **(4, 22)**  
['P300\_25', 'P1000\_30', 'seed\_nb'] **(3, 25)**  
['P300\_25', 'P1000\_30', 'seed\_yield'] **(3, 24)**  
['P300\_25', 'P1000\_30', 'PMG'] **(3, 24)**  
['P1000\_30', 'PMG', 'lipids'] **(3, 24)**  
['climate', 'genotype', 'PMG'] **(3, 22)**  
['P600\_25', 'P1000\_30', 'seed\_yield'] **(3, 22)**  
['P600\_25', 'P1000\_30', 'proteins'] **(3, 22)**