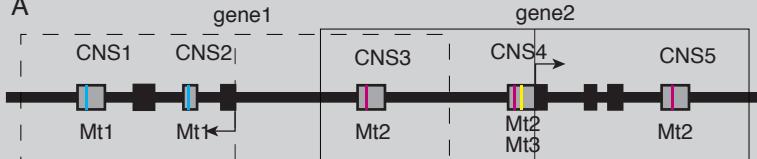
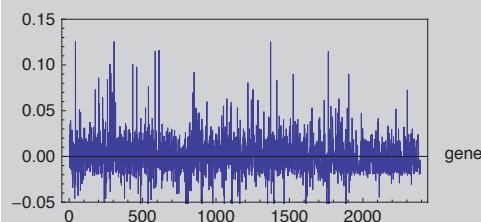


A**B**

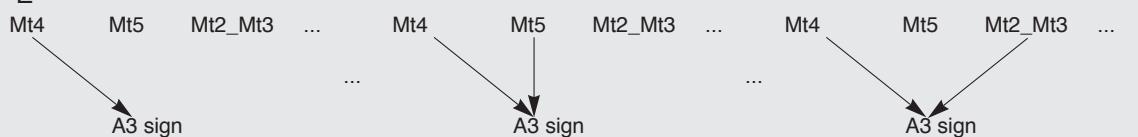
Eigenarray A3

**C**

| | Mt1 | Mt2 | Mt3 | ... | |
|-------|-------|---------|-----|---------------------------------|-------------------------|
| gene1 | {1,2} | {3} | 0 | ... | |
| gene2 | 0 | {3,4,5} | {4} | ... | |
| ... | ... | ... | ... | ... | |
| Mt1 | Mt2 | Mt3 | ... | Mt1_Mt2 Mt1_Mt3 ... Mt2_Mt3 ... | |
| gene1 | 1 | 1 | 0 | ... | 0 0 ... 0 0 ... 0 0 ... |
| gene2 | 0 | 1 | 1 | ... | 0 0 ... 0 0 ... 1 0 ... |
| ... | ... | ... | ... | ... | ... |

D

| | A1 | A2 | A3 | ... |
|-------|-----|-----|-----|-----|
| gene1 | + | - | + | ... |
| gene2 | + | + | - | - |
| ... | ... | ... | ... | ... |

E**F**

| BN score | q-value | Set of Comp. Motifs |
|-----------------------|---------|---------------------|
| 4.3107×10^8 | <0.001 | Mt4 Mt5 |
| 4.17003×10^8 | <0.001 | Mt5 Mt61 |
| 1.5724×10^8 | <0.001 | Mt4_Mt61 Mt5 |
| ... | ... | ... |

| motifs | | A3 sign | | number of genes |
|--------|-----|---------|------|-----------------|
| Mt4 | Mt5 | - | + | |
| 0 | 1 | 0.5 | 0.5 | 96 |
| 1 | 0 | 0.48 | 0.52 | 381 |
| 0 | 0 | 0.57 | 0.43 | 328 |
| 1 | 1 | 0.21 | 0.79 | 234 |
| All | | 0.45 | 0.55 | 1039 |