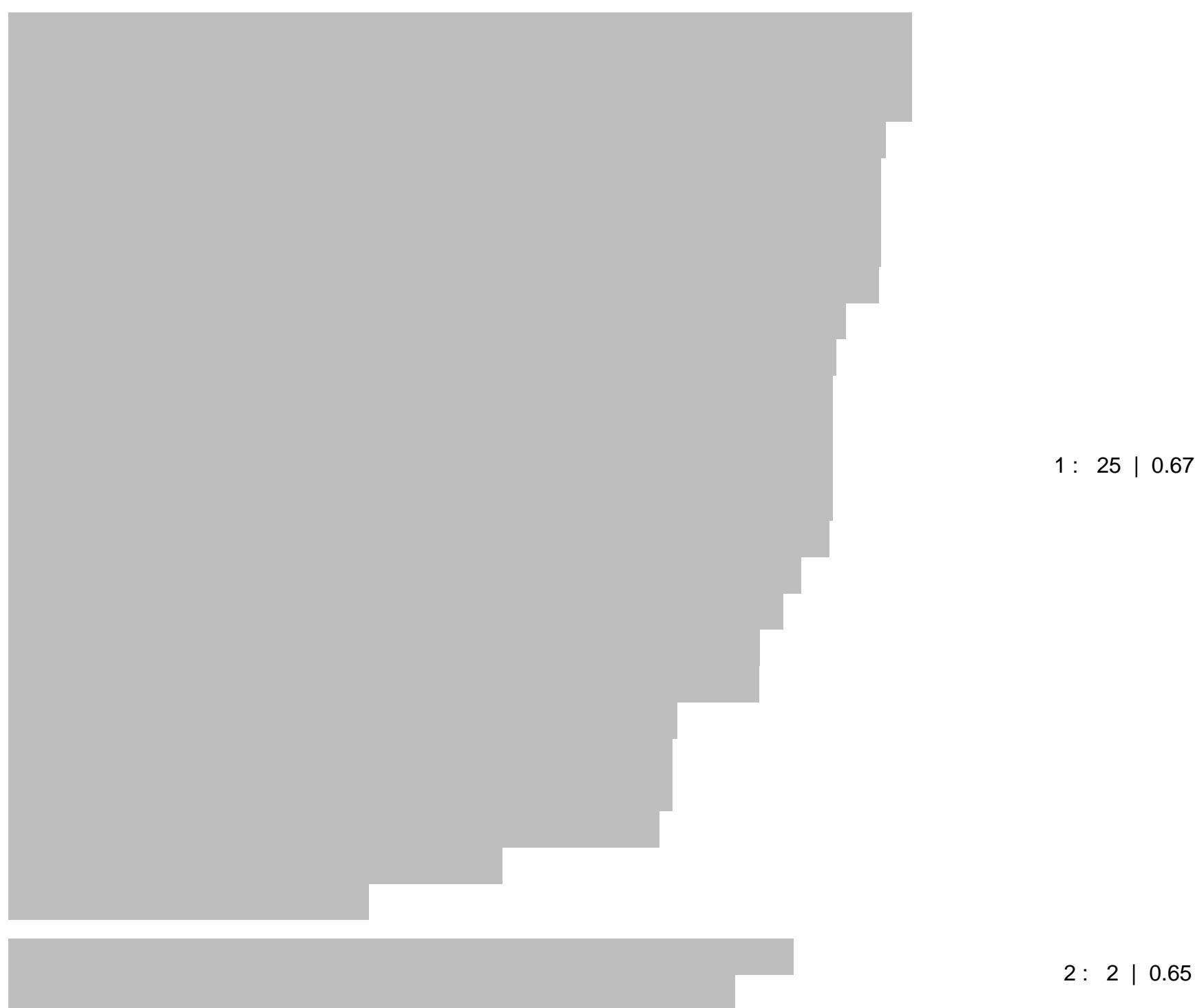


## Silhouette – Genotype Form linear

n = 27

2 clusters  $C_j$   
 $j : n_j | \text{ave}_{i \in C_j} s_i$

CV\_t  
gPi  
CEV  
SI  
PPi  
Plmd  
PILSM  
PFI  
EVS  
PIR  
ESPI  
PQ  
PR  
NRW  
ESPII  
D\_slo  
APC  
ESP  
PPF  
RC  
RDPI  
RPI  
RNN  
RN  
RTT  
RSI  
PSI



0.0 0.2 0.4 0.6 0.8 1.0

Silhouette width  $s_i$

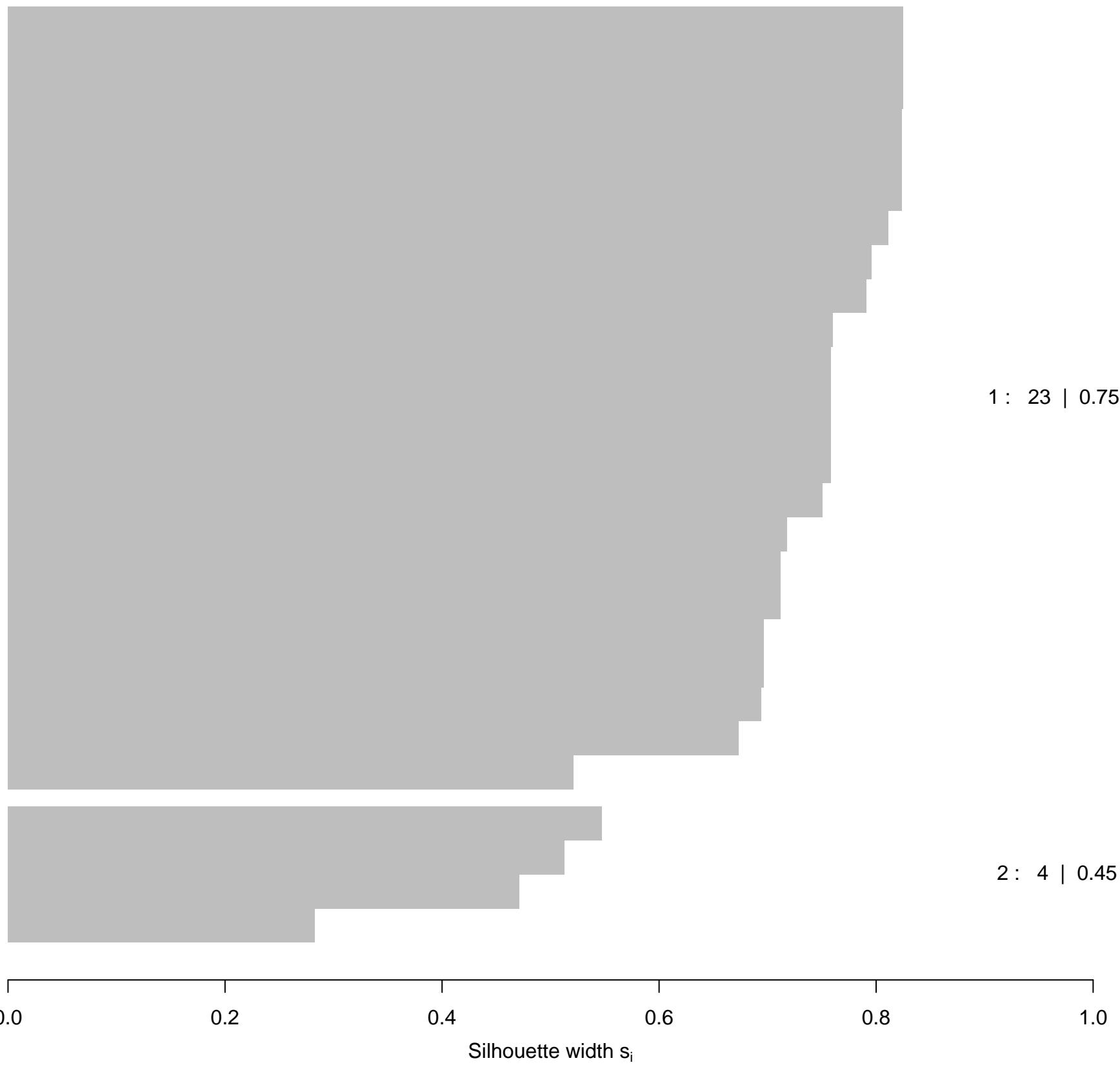
Average silhouette width : 0.67

## Silhouette – Genotype Form gaussian

n = 27

2 clusters  $C_j$   
 $j : n_j | \text{ave}_{i \in C_j} s_i$

PPi  
Plmd  
PILSM  
CV\_t  
gPi  
CEV  
PFI  
PIR  
SI  
ESP  
ESPI  
PQ  
PR  
NRW  
EVS  
D\_slo  
ESPII  
APC  
RDPI  
RPI  
RC  
PPF  
RNN  
PSI  
RSI  
RN  
RTT



Average silhouette width : 0.71

## Silhouette – Genotype Form sinusoidal

n = 27

2 clusters  $C_j$   
 $j : n_j | \text{ave}_{i \in C_j} s_i$

PPi

Plmd

PILSM

CV\_t

gPi

CEV

SI

PIR

PFI

PPF

RDPI

RPI

EVS

ESPI

PQ

PR

NRW

RC

ESP

D\_slo

ESPII

RNN

APC

PSI

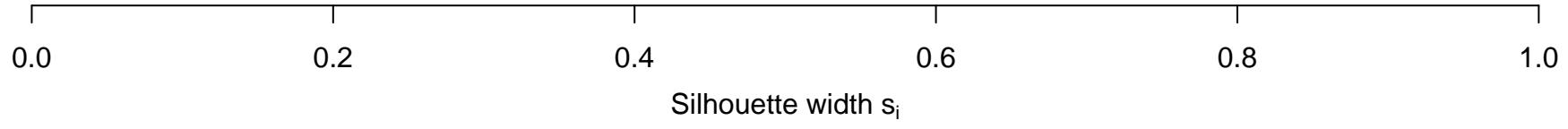
RN

RSI

RTT

1 : 23 | 0.82

2 : 4 | 0.74



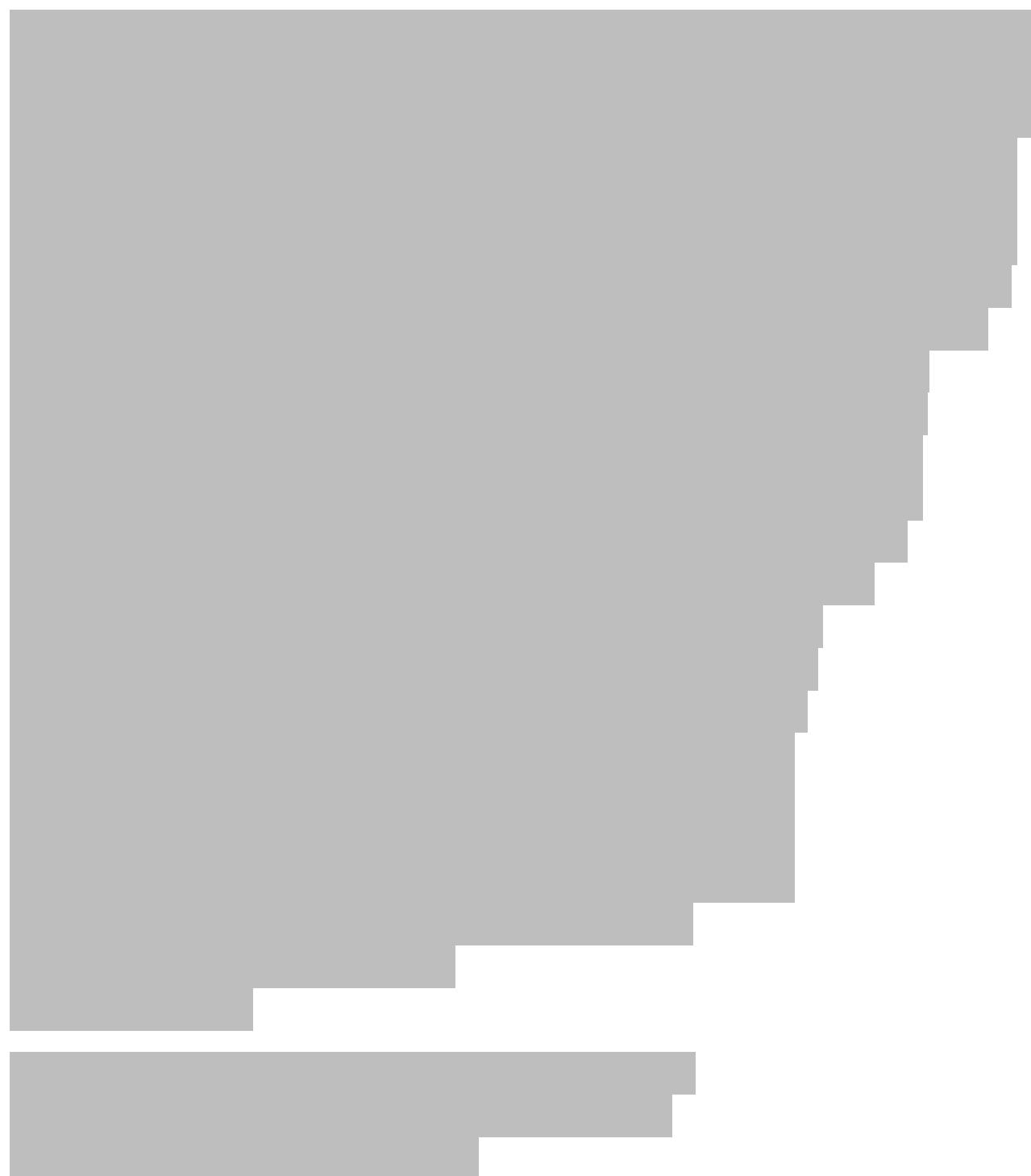
Average silhouette width : 0.81

## Silhouette – Genotype Form wave

n = 27

2 clusters  $C_j$   
 $j : n_j | \text{ave}_{i \in C_j} s_i$

CV\_t  
gPi  
CEV  
PPI  
Plmd  
PILSM  
ESP  
SI  
RC  
PFI  
RDPI  
RPI  
PIR  
PPF  
EVS  
ESPII  
D\_slo  
ESPI  
PQ  
PR  
NRW  
APC  
RNN  
PSI  
RTR  
RSI  
RN



1 : 24 | 0.63

2 : 3 | 0.45

0.0 0.2 0.4 0.6 0.8 1.0

Silhouette width  $s_i$

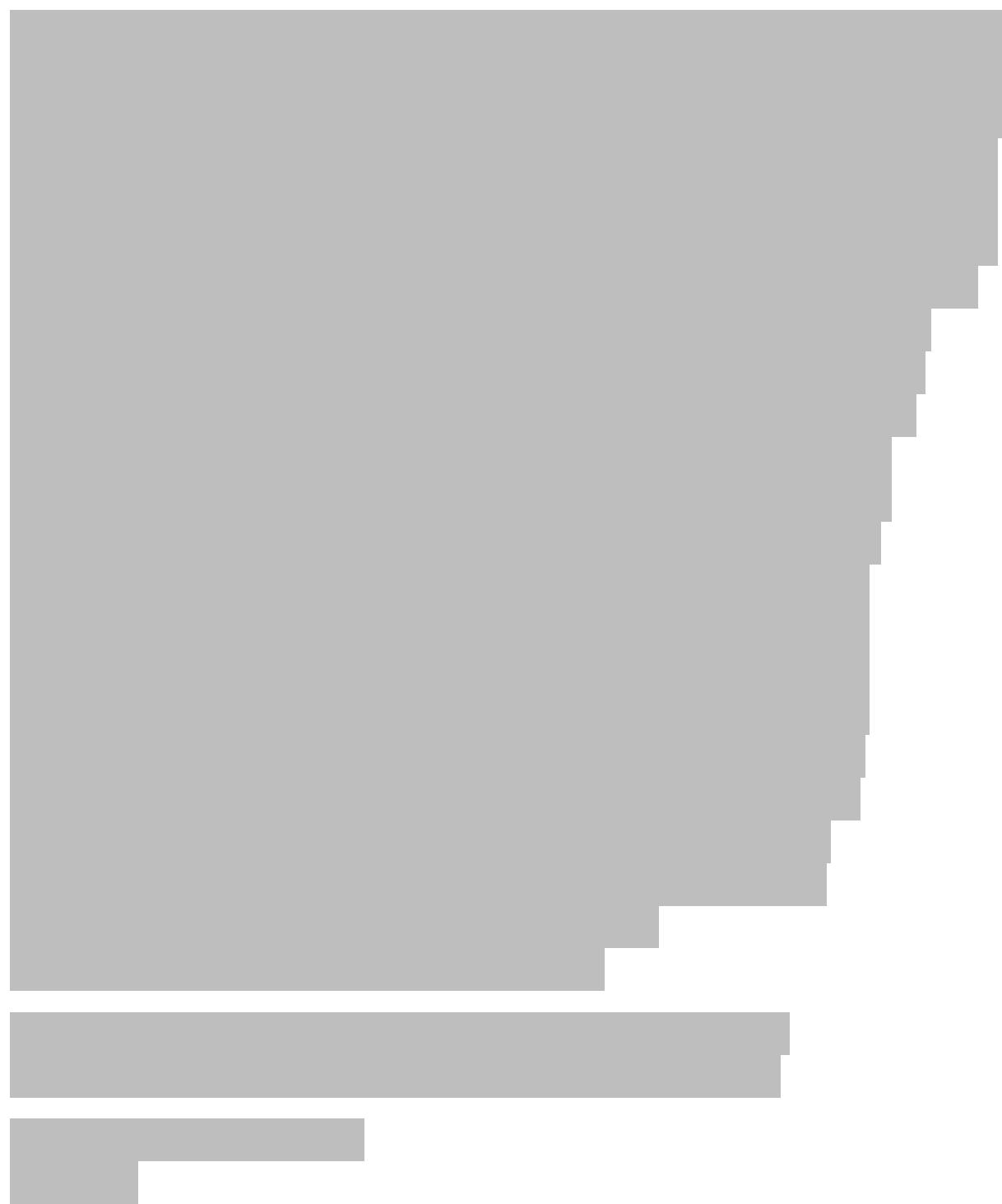
Average silhouette width : 0.61

## Silhouette – Combined Analysis

n = 27

3 clusters  $C_j$   
 $j : n_j | \text{ave}_{i \in C_j} s_i$

PPi  
Plmd  
PILSM  
CV\_t  
gPi  
CEV  
SI  
ESP  
PIR  
PFI  
RDPI  
RPI  
RC  
ESPI  
PQ  
PR  
NRW  
EVS  
D\_slo  
ESPII  
PPF  
APC  
RNN



1 : 23 | 0.64

2 : 2 | 0.56

3 : 2 | 0.17

0.0 0.2 0.4 0.6 0.8 1.0

Silhouette width  $s_i$

Average silhouette width : 0.6