## Supplementary Materials for

# Discovery of ROH diplotype clusters and their associations with diseases in UK Biobank

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#### Algorithm 1 Find all blocks of matches while maximizing the number of haplotypes

```
procedure MAXIMAIZE HAPLOTYPES (x_k, t, a_k, d_k, L, c)
    V = [], Q = []
    for r = 0 to t do
       m.append([])
        pq.append(k+1)
        V.append([])
        Q.append([])
    end for
    io ← 0
    for i = 0 to M - 1 do
       if d_k[i] > k - L + 1 then
            changed← false
            \mathbf{for}\,p = 0 \text{ to } t \text{ do}
               for q = p + 1 to t do
                   if len(m[p]) > 0 and len(m[q] > 0) then
                       changed ← true
                        break
                    end if
                end for
            end for
            if changed and i - i0 >= c then
               blockMatches = []
               d_{min} \leftarrow 0
                blockMatches.append[a_k[io]]
               for i_b = i_0 + 1 to i do
                    blockMatches.append([ak[ib]])
                   if (dk[i_b] > d_{min}) then
                        d_{min} = div[i_b]
                    end if
                end for
               report dmin and blockM atches
            end if
            i0 \leftarrow i, m = []
            \mathbf{for} \, r = 0 \, \mathbf{to} \, t \, \mathbf{do}
               m.append([])
            end for
        end if
        updateV_Q(x_k[l], i, t, a_k, d_k, pq, V, Q)
    a_{k+1} \leftarrow concatenate[Vo, V_1, ...V_t]
    d_{k+1} concatenate[Q0, Q1, ..Qt]
end procedure
```

### Algorithm 2 Update intermediate variables for cPBWT

```
procedure UPDATEV_Q(allele, i, t, a_k, d_k, pq, V, Q)

for p = 0 to t do

if d_k[i] > pq[p] then

pq[p] \leftarrow d_k[i]

end if

end for

V [allele].append(a_k[i])

for p = 0 to t do

if allele == p then

Q[allele].append(pq[allele])

pq[allele] \leftarrow 0

mab[p].append(a_k[i])

end if

end for

end procedure
```

#### **Algorithm 3** Find all blocks of matches larger than L while maximizing the length of match

```
procedure Maximaize haplotypes(x_k, t, a_k, d_k, L, c)
             V = [], Q = []

for r = 0 to t do

m.append([])

pq.append([])

V.append([])
                    Q.append([])
              end for
              i_0 \leftarrow 0
              for i = 0 to M - 1 do
                   if d_k[i] > k - L + 1 then
                          changed \leftarrow f \text{ also}
changed \leftarrow f \text{ also}
for p = 0 \text{ to } t \text{ do}
for q = p + 1 \text{ to } t \text{ do}
if len(m[p]) > 0 \text{ and } len(m[q] > 0) \text{ then}
changed \leftarrow true
                                              break
                                 end if
end for
                           end for
                          if changed and i - i_0 >= c then
blockMatches = [], d_{min} \leftarrow 0, f_a \leftarrow a_k[i_0], f_d \leftarrow d_k[i_0 + 1]
T_d[f_a].append(f_d), all_d.append(f_d)
                                 for i_a = i_0 + 1 + 0 i do blockMatches.append([a_k[i_b]]) if (d[i_a] > d_{min}) then d_{min} = d_k[i_b] end if
                                 f_a \leftarrow x_k[a_k[i_a]], \ T_d[f_a]. \ append(d_{min}), \ all_d. \ append(d_{min}) end for
                                 for r = 0 to t do

if len[T_d[r]] >= c and quick \ select(T_d[r], c - 1) <= quick \ select(all_d, c - 1) then
                                              skip
                                              i_a \leftarrow i_0

if f_d \le quick select(all_d, c - 1)) then
                                                     block Matches.append ([a_k[i_0]]
                                                     for i_a = i_0 + 1 to i do

if d_k[i_a] \le quick\_select(all_a, c - 1) then

blockMatches.append(a_k[i])
                                                           end if
                                                     end for
                                                    d_r = quick \ select(all_d, c - 1)
report blockMatches and d_r
                                              end if
                                              i_0 \leftarrow i, m = []

for r = 0 to t do
                                              \begin{array}{c} \text{m.append}([]) \\ \textbf{end for} \end{array}
                                        end if
                                        updateVQ(x_k[l], i, t, a_k, d_k, pq, V, Q)
                                  end for
                                 end for a_{k+1} \leftarrow concatenate[V_0, V_1, ...V_t]

d_{k+1} \leftarrow concatenate[Q_0, Q_1, ...Q_t]
```