Figure 2. Case and Control Selection.

HIV-1 Infection
Subtype B (USA, Trinidad and Tobago)
Subtype C (Sub-Saharan Africa)

Target Population
\( n = 12,590 \)
Sequences from Los Alamos HIV Database

Cases
\( n = 133 \)

Assessed for Eligibility
\( n = 766 \)

Participants
\( n = 133 \)

Eligible Controls
\( n = 133 \)

Excluded \( n = 633 \)
- Incomplete sequences \( n = 303 \)
- From acute infection \( n = 168 \)
- Isolated from brain/CSF \( n = 12 \)
- Isolated from genital tract \( n = 2 \)
- Isolated from vaccine trial \( n = 1 \)
- Isolated from chimp \( n = 1 \)
- Virus created in lab \( n = 3 \)
- Virus manipulated \textit{in vitro} \( n = 21 \)