

Keywords employed to pinpoint sequences from the built environment:

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be_list = ["HEPA", "air", "attic", "building", "built", "carpet",  
          "cellar", "cleaning", "cleanroom", "construction", "door",  
          "dust", "filter", "floor", "gypsum", "home", "hospital",  
          "house", "indoor", "room", "space", "theatre", "wall",  
          "ventilation"]
```

These were matched against

1. The title of the underlying scientific study in the INSDC
2. The INSD source field
3. The INSD tissue type field
4. UNITE sequence source (either "Air" or "Dust fungal DNA")