The graph illustrates the normalized expression levels of various genes in NSCLC and Tuberculo pleurisy samples. The x-axis represents the number of undetected samples, and the y-axis shows the normalized expression levels. The y-axis range is from 1E-7 to 1, with markers for 1E-3, 1E-4, 1E-5, and 1E-6.

- **Gene Expression Information**:
  - **LunX**: P < 0.001
  - **CK19**: P = 0.066
  - **CEA**: P = 0.074
  - **VEGF-C**: P = 0.054
  - **hnRNP A2/B1**: P = 0.613

- **Marker Definitions**:
  - Black dots represent NSCLC samples (n=14).
  - Open circles represent Tuberculo pleurisy samples (n=14).

The data suggests significant differences in gene expression between the two conditions, with LunX and CEA showing the most pronounced differences.