LaTeX Author Guidelines for 8.5×11-Inch Proceedings Manuscripts

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0.1. Algorithm

Let E be some experiment from a research paper with N participants . Then we have that $E = \{P_1, P_2, P_3, ..., P_N\}$. Each participant in the experiment P_i has a set of l microbial samples taken from a particular body site, and each sample m has a time when it was taken t. Then we have that $S^t(P_i) = \{m_{S_1}, m_{S_2}, ..., m_{S_l}\}$ where each S_j is a sample from participant P_i, and $i > 0$, $j > 0$, and $l > 0$. Then each microbial sample m_{S_j} is a set of x operational taxonomic units or bugs such that we have $m_{S_j} = \{b_1, b_2, ..., b_x\}$ where $x > 0$.

Assume an attacker has k samples taken from a single individual at a time t referred to as mr.duck smith where k greater than or equal to 1. Then $A = \{a_1, a_2, a_3, ..., a_k\}$ is the set of the samples from mr. duck smith that the attacker has collected.

Then let P be all the participants in experiment E, S be all the samples in experiment E, and O be all the possible operational taxonomic units (OTU) . Let N be the number of participants in experiment E, M be the number of samples in experiment E, and L be the number of OTUs in O.

Then the following procedures define an algorithm for determining if mr. duck smith is a participant in experiment E :

Algorithm 1 Probability Based Microbial Signature Algorithm

```
1: procedure MAKESIGNATURE(P_i,S,O,percent)
2:     i ← 0
3:     j ← 0
4:     count ← 0
5:     sig ← []
6:     for i = 1 → length(O) do
7:         for j = 1 → length(S) do
8:             sample ← S[j]
9:             if O[i] ∈ sample then
10:                count ← count + 1
11:            totalPercent ← count \ length(S)
12:            if totalPercent ≥ percent then
13:                sig ← sig.append(O[i], 1)
14:            else
15:                sig ← sig.append(O[i], 0)
16:            count ← 0
17: procedure REMOVEOTUS(P, Sigs, O, percent)
18:    i ← 0
19:    j ← 0
20:    k ← 0
21:    count ← 0
22:    OTUs ← []
23:    for i = 1 → length(O) do
24:        for j = 1 → length(P) do
25:            if O[i] ∈ Sigs(P_j) then
26:                count ← count + 1
27:            totalPercent ← count \ length(S)
28:            if totalPercent ≥ percent then
29:                OTUs ← OTUS.append(O[i])
30:        count ← 0
31:    for k = 1 → length(Sigs) do
32:        Sigs[k] ← Sigs[k].remove(OTUs)
33: procedure MATCHSIGNATURE(s_1,s_2)
34:    i ← 0
35:    count ← 0
36:    for i = 1 → length(S_1) do
37:        if S_1[i]xorS_2[i] then
38:            continue
39:        else
40:            count ← count + 1
41:    return count
42: procedure MATCHSIGNATURE(Sigs,aSig)
43:    i ← 0
44:    samps ← []
45:    for i = 1 → length(Sigs) do
46:        value ← MatchSignature(Sigs[i], aSig])
47:        samps ← samps.append((Sigs[i]), value)
48:    sort samps in ascending order return samps
```