

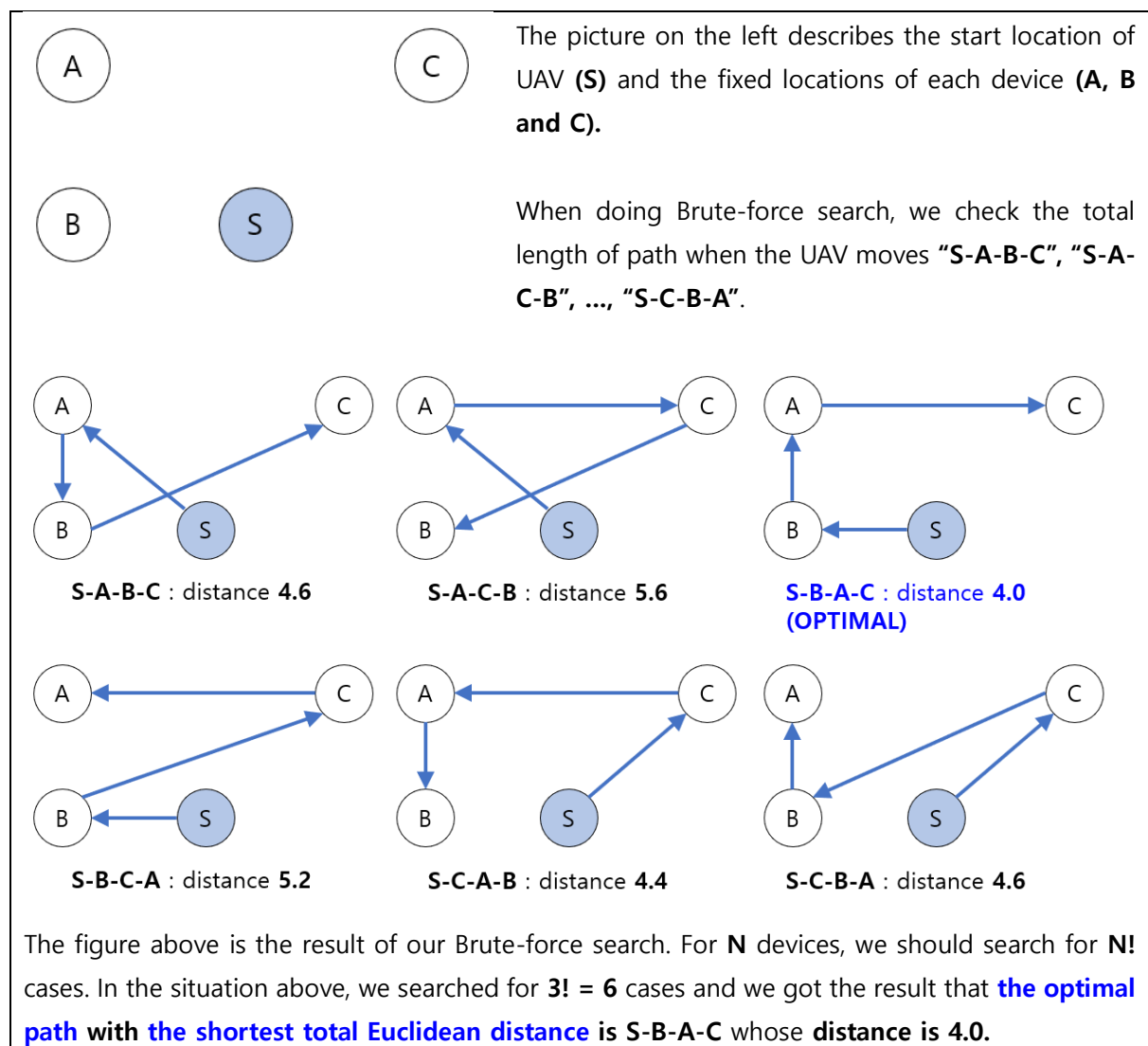
FINAL DOCUMENTATION FOR WPCN-UAV PAPER

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1. newGeneticDeviceVisitAl.py

1-1. **Goal:** For each cluster, to determine whether the brute-force search of the optimal path for the movement of UAV

1-2. **Brute-force search of optimal path:**



1-3. Swap-basic search:

Swap-basic search is the method to find the shortest total Euclidean distance with swapping the device visit sequence. **(suppose that the number of device is 8)**

With start location of UAV S , device $d1, d2, \dots, d8$ where **CURRENT (updated)** visit sequence is $S \rightarrow d1 \rightarrow d2 \rightarrow \dots \rightarrow d8$

While True: // until convergence

For each **case** in $\{d1, d2, d3\}, \{d1, d2, d3, d4\}, \dots, \{d4, d5, d6, d7\}, \{d5, d6, d7, d8\}, \{d6, d7, d8\}$ // total $(8 - 1 = 7)$ cases

If **$d1$ in case** and $\text{len}(\text{case}) == 3$: // include start

swapCondition = $(S \rightarrow d2 \rightarrow d1 \rightarrow d3)$ is shorter than $(S \rightarrow d1 \rightarrow d2 \rightarrow d3)$

if swapCondition is True:

$(d1, d2) = (d2, d1)$

else if **$d8$ in case** and $\text{len}(\text{case}) == 3$: // include end

swapCondition = $(d6 \rightarrow d8 \rightarrow d7)$ is shorter than $(d6 \rightarrow d7 \rightarrow d8)$

if swapCondition is True:

$(d7, d8) = (d8, d7)$

else: // other cases

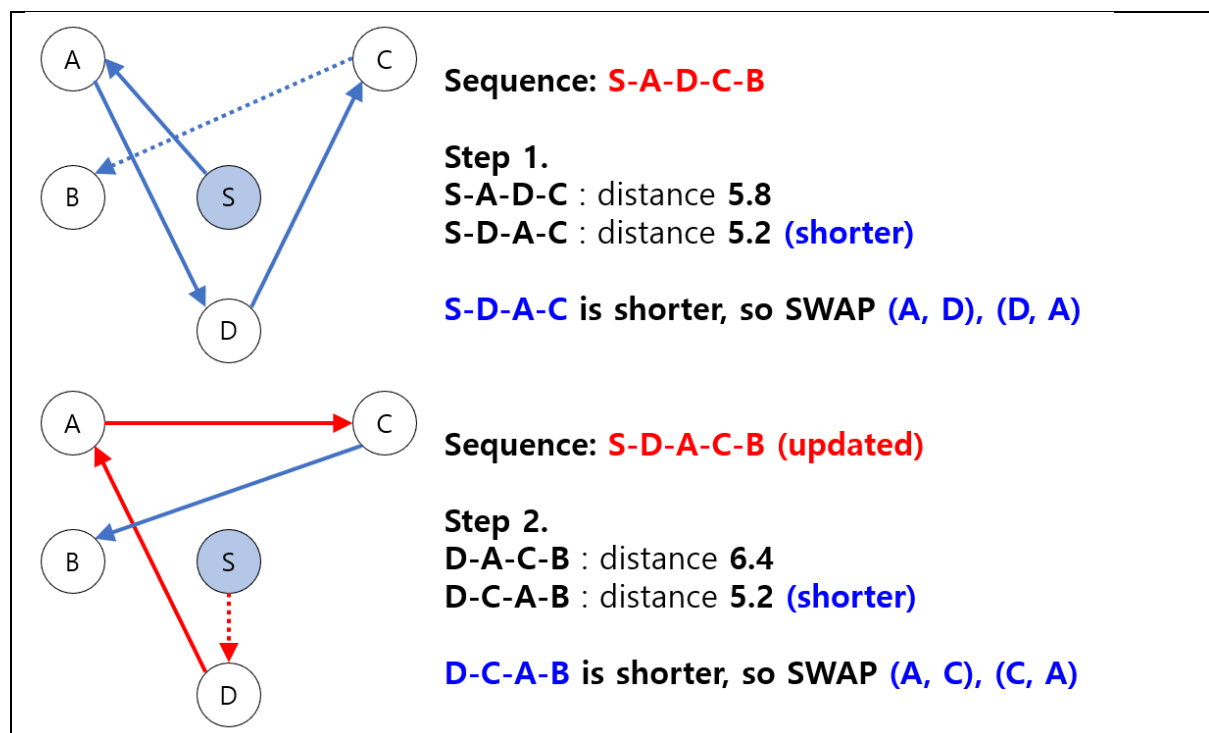
swapCondition = $(\text{case}[0] \rightarrow \text{case}[2] \rightarrow \text{case}[1] \rightarrow \text{case}[3])$ is shorter than $(\text{case}[0] \rightarrow \text{case}[1] \rightarrow \text{case}[2] \rightarrow \text{case}[3])$

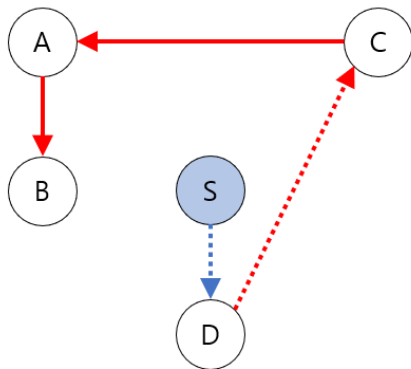
if swapCondition is True:

$(\text{case}[1], \text{case}[2]) = (\text{case}[2], \text{case}[1])$

Terminate if **do not updated** // terminate when reached convergence

For example,





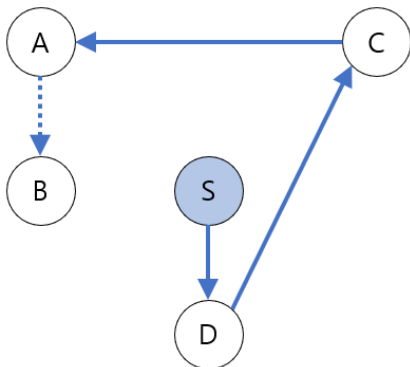
Sequence: **S-D-C-A-B (updated)**

Step 3.

C-A-B : distance **3.0 (shorter)**

C-B-A : distance **3.2**

C-A-B is shorter, so **DO NOT SWAP**



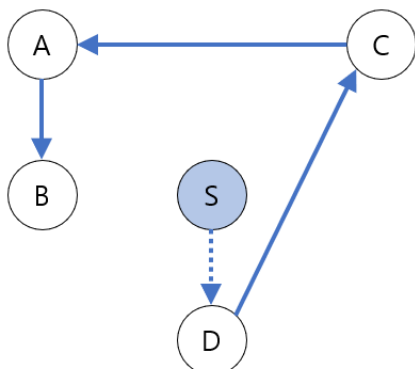
Sequence: **S-D-C-A-B**

Step 4.

S-D-C-A : distance **5.2 (shorter)**

S-C-D-A : distance **5.8**

S-D-C-A is shorter, so **DO NOT SWAP**



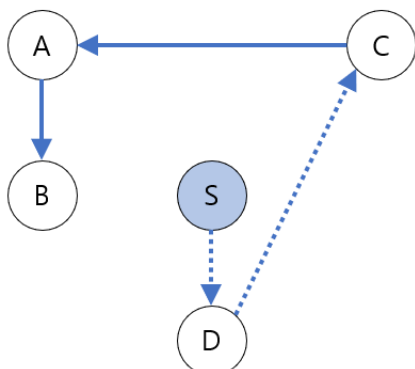
Sequence: **S-D-C-A-B**

Step 5.

D-C-A-B : distance **5.2 (shorter)**

D-A-C-B : distance **6.4**

D-C-A-B is shorter, so **DO NOT SWAP**



Sequence: **S-D-C-A-B**

Step 6.

C-A-B : distance **3.0 (shorter)**

C-B-A : distance **3.2**

C-A-B is shorter, so **DO NOT SWAP**
Terminate because **NO UPDATE**

1st iteration : Step 1 ~ Step 3

2nd iteration : Step 4 ~ Step 6

1-4. Deep Learning Approach

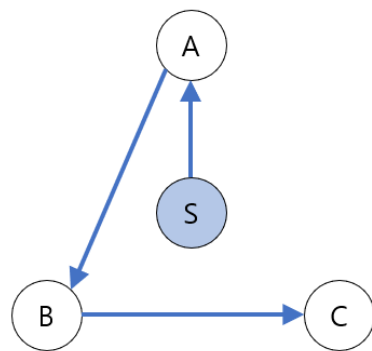
Use Deep Learning to **determine whether the Brute-force search is needed**:

Original data	Movement S->d1->d2->...->d8
Input data	<p>For each segment of the movement S->d1, d1->d2, ..., d7->d8,</p> <ol style="list-style-type: none"> x distance of the movement y distance of the movement <p>For each segment of the movement S->d1, d1->d2, ..., d7->d8,</p> <ol style="list-style-type: none"> sin(A) where A is the angle of the movement cos(A) where A is the angle of the movement Euclidean distance of the movement
Output data	<p style="text-align: center;">$\min\left(1.0, \log_{1.75} \frac{D_{SB}}{D_{BF}}\right) \rightarrow \text{always between } 0.0 \text{ and } 1.0$</p> <p>Where D_{SB} is the total length of the movement path after applying Swap-Basic method, and D_{BF} is the total length of the movement path of the optimal movement derived by Brute-force search</p> <p>(NOTE: $D_{SB} \geq D_{BF}$ is always true because D_{BF} is the optimal distance)</p> <div style="display: flex; align-items: flex-start;"> <div style="flex: 1;"> </div> <div style="flex: 1;"> <p>Sequence: S-D-C-A-B</p> <p>Step 6. C-A-B : distance 3.0 (shorter) C-B-A : distance 3.2</p> <p>C-A-B is shorter, so DO NOT SWAP Terminate because NO UPDATE</p> </div> </div> <p>In the situation of 1-3 (the picture above), the movement path of S->D->C->A->B is derived by Swap-Basic method and whose total distance is 6.2, so $D_{SB} = 6.2$</p> <div style="display: flex; align-items: flex-start;"> <div style="flex: 1;"> </div> <div style="flex: 1;"> <p>In the situation of 1-2, (the picture on the left side) the movement path of S->B->A->C is optimal (derived by Brute-force method) and whose total distance is 4.0, so $D_{BF} = 4.0$</p> <p>S-B-A-C : distance 4.0 (OPTIMAL)</p> </div> </div> <p>NOTE: The two cases above are DIFFERENT (so, DO NOT think of 6.2/4.0)</p>

Suppose that **the start location of the UAV is the CENTROID of all the devices** and **the number of devices is 8**

Main idea (important): Brute-force search is needed if D_{SB}/D_{BF} value is large enough

Example about the input data:



movement path: **S->A->B->C**

location of the start location of UAV (**S**) and each device (**A, B and C**) is,

- ➔ S (x=**0.0**, y=**0.0**)
- ➔ A (x=**0.0**, y=**+1.0**)
- ➔ B (x=**-0.866**, y=**-0.5**)
- ➔ C (x=**+0.866**, y=**-0.5**)

For each segment **S->A, A->B and B->C**,

	S->A	A->B	B->C
X distance	X1 = 0.0	X2 = -0.866	X3 = +1.732
Y distance	Y1 = +1.0	Y2 = -1.5	Y3 = 0.0
Angle A	90 degree	240 degree	0 degree
Sin(A)	Sin1 = +1.0	Sin2 = -0.866	Sin3 = 0.0
Cos(A)	Cos1 = 0.0	Cos2 = -0.5	Cos3 = +1.0
Euclidean distance	Dist1 = 1.0	Dist2 = 1.732	Dist3 = 1.732

The form of the input data is,

(X1, Y1, X2, Y2, X3, Y3, sin1, cos1, dist1, sin2, cos2, dist2, sin3, cos3, dist3)

So, the input data for this case is,

(0.0, +1.0, -0.866, -1.5, +1.732, 0.0, +1.0, 0.0, +1.0, -0.866, -0.5, +1.732, 0.0, +1.0, +1.732)

1-5. Model Genetation using the idea of 1-4

1. Create **16,000** cases of the clusters (refer to variable '**times**' of **newGeneticDeviceVisitAI.py**)

➔ The number of devices in the cluster is as the table below

2. Do Deep Learning for these cases using **12,960 (81%) training data, 1,440 (9%) validation data and 1,600 (10%) test data**
3. The model is generated based on the deep learning result.

```

if __name__ == '__main__':
    # numpy setting
    np.set_printoptions(edgeitems=20, linewidth=200)

    input_data = []
    output_data = []
    times = 16000
    deviceCountList = []
  
```

The number of devices: **(probability)**

4 devices	5	6	7	8 devices
15%	22%	26%	22%	15%

2. throughputTest_NewGenetic.py

2-1. Configuration

In **settings.txt**, you can modify the variables below:

	description	Default value
width	Width in meter	50
height	Height in meter	50
L	The number of clusters	5
devices	The total number of devices in the all clusters	30
N	Flight period (the number of time slots)	30
iters	The number of iterations	1,200
clusteringAtLeast	cluster contains devices at least $\text{int}(\text{this number} * \text{average})$	0.67
clusteringAtMost	cluster contains devices at most $\text{int}(\text{this number} * \text{average})$	1.34

The total number of cases (clusters) to test: **L * iters (default 5 * 1,200 = 6,000)**

In the setting above, the average number of device for each cluster is **(device / L) = 30 / 5 = 6**, so the cluster contains at least **$\text{int}(0.67 * 6) = 4$ devices**, and at most **$\text{int}(1.34 * 6) = 8$ devices**. It is similar to the distribution of the number of the devices in 1-5.

2-2. Mode Settings


Static (static_swapOnly)	Same as the static strategy from Tang et al → Tang J, Song J, Ou Jea. Minimum Throughput Maximization for Multi-UAV Enabled WPCN: A Deep Reinforcement Learning Method. IEEE Access 2020.
Train swap only (train_swapOnly)	Find the device-visit sequence (movement path) using Swap-basic search , and then visit devices using this sequence. Compute the common throughput based on the sequence.
Train gen visit (train_genVisit)	Find the device-visit sequence (movement path) using Swap-basic search , and then input the movement path to the model generated in 1-5 . If the output value of the model is at least (threshold = 0.5) , then find the device-visit sequence using Brute-force search and then visit devices using this sequence. Otherwise, visit devices using the sequence derived by Swap-basic search . Compute the common throughput based on the sequence of device-visit .


In () is **the prefix of the trajectory image file name**. (**static_swapOnly_trajectory_iter0000.png** for example)


2-3. The number of iterations for each mode

mode	The number of iterations	Total generated clusters when iters=1,200, L=5
Static	iters // 2	$(1,200 // 2) * 5 = 3,000$
Train swap only	iters	$1,200 * 5 = 6,000$
Train gen visit	iters	$1,200 * 5 = 6,000$

2-4. interpretation of the result

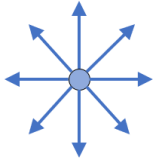
Static mode
 minThroughputList_static2208010048_swapOnly_iter_1200_L_0005_devs_0030_N_0030.txt - Windows 메모장 파일(F) 편집(E) 서식(O) 보기(V) 도움말(H) mean: 0.01836036916346323, std: 0.007857989538148612, nonzero: 3000 Time: 2022.08.01 00:48 Mode: Static (refer to static2208010048_swapOnly) Configurations: iters= 1200 , L= 5 , devices= 30 , N= 30 Average value of the common throughput: 0.018360 Standard deviation of the common throughput: 0.007858 The total number of clusters whose common throughput is non-zero: 3,000 / 3,000 (100.0%)

Train swap only
 minThroughputList_train2208010048_swapOnly_iter_1200_L_0005_devs_0030_N_0030.txt - Windows 메모장 파일(F) 편집(E) 서식(O) 보기(V) 도움말(H) mean: 0.02500243346203403, std: 0.011188922783291363, nonzero: 5994 Mode: Train swap only (refer to train2208010048_swapOnly) Average value, standard deviation of the common throughput: 0.025002, 0.011189 The total number of clusters whose common throughput is non-zero: 5,994 / 6,000 (99.9%)

Train gen visit
 minThroughputList_train2208010048_genVisit_iter_1200_L_0005_devs_0030_N_0030.txt - Windows 메모장 파일(F) 편집(E) 서식(O) 보기(V) 도움말(H) mean: 0.026219649894172997, std: 0.010857653190466882, nonzero: 5999 Mode: Train gen visit (refer to train2208010048_genVisit) Average value, standard deviation of the common throughput: 0.026220, 0.010858 The total number of clusters whose common throughput is non-zero: 5,999 / 6,000 (99.98%)

3. Actual movement of UAV

3-1. actual movement in each segment



Unit movement is the movement of the UAV in each time slot. Its distance is always **5.0m** and the degree of direction is always one of **0-degree, 45-degree, 90-degree, ..., and 315-degree**.

For each unit movement of the movement path **from device A to device B**,

1. Find the direction of the next device
2. Move toward the direction
3. Stop until **moving toward any of the 8 directions increments the distance** between UAV and the device

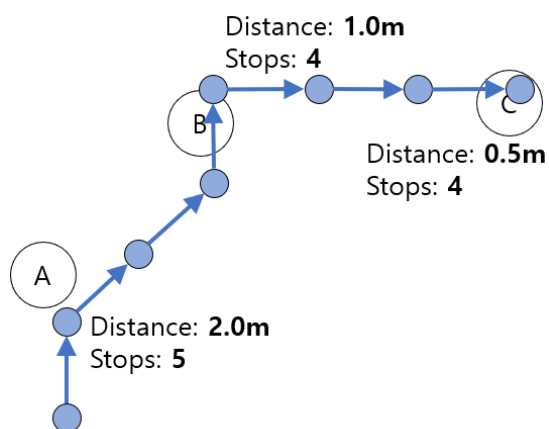
3-2. communication with devices

1. If there are at least 1 device **within 3.125m** of the current UAV location, **communicate with these devices in sequence**.
2. Otherwise, communicate with **the most nearby** device.

3-3. final decision of the movement (with devices **d1, d2, ..., d8**)

1. Compute the actual movement of **S->d1, d1->d2, ..., d7->d8**, and assign time slots to the movement.
2. Assign each remaining time slots to the device (**stop UAV for these time slots**), based on **the distance between the UAV (in proportion to $(distance)^{0.002}$)** and the device when **the UAV is closest to the device**.


For example, if there are **20 time slots** in total and device **A, B and C**,



1. assign **1 time slot** to **S->A**, **3 time slots** to **A->B**, and **3 time slots** to **B->C**, according to **3-1** for each segment **S->A, A->B and B->C**.
2. Assign remaining 13 time slots to stopping at the closest device.
 - ➔ The closest distance between UAV and each device is **2.0m, 1.0m, 0.5m** for device **A, B and C**.
 - ➔ So, assign **5 time slots** to the farthest device **A**, and **4 time slots** for other devices.

4. Running

4-1. run newGeneticDeviceVisitAI.py first

 Genetic_Visit_AI_model


2022-08-01 오전 12:47 파일 폴더

If you already have Genetic_Visit_AI_model directory with the model, you can skip this stage.

```
D:\WPNUAV>python newGeneticDeviceVisitAI.py
D:\Python37\lib\site-packages\tensorflow_core\python\pywrap_tensorflow_internal.pyd
0
input data (swap) : [ 1.7474  0.4497 -1.3823  1.304  -1.2989 -2.3533 -0.0068 -0.3983  0.
input data (angle) : [ 0.2492  0.9684  1.8043  0.6862 -0.7274  1.9003 -0.8755 -0.4832  2.
,      0.      0.      0.      0.      ]
dist      : 1.7676 1.4795
0 [1, 0, 2, 3, 4]
1 [4, 3, 2, 0, 1]
```

Otherwise, run **python newGeneticDeviceVisitAI.py.**

4-2. run throughputTest_NewGenetic.py

 명령 프롬프트

```
D:\WPNUAV>python throughputTest_NewGenetic.py
D:\Python37\lib\site-packages\tensorflow_core\python\pywrap_tensorflow_internal.pyd
```

Run the command **python throughputTest_NewGenetic.py.**