

In [30]:

```
# Import Libraries
from matplotlib import pyplot as plt
import numpy as np
import pandas as pd
file=pd.read_csv('C:/Users/wit5/Downloads/skin_cancer.csv')
file
```

Out[30]:

|       | lesion_id   | image_id     | dx    | dx_type | age  | sex    | localization |
|-------|-------------|--------------|-------|---------|------|--------|--------------|
| 0     | HAM_0000118 | ISIC_0027419 | bkl   | histo   | 80.0 | male   | scalp        |
| 1     | HAM_0000118 | ISIC_0025030 | bkl   | histo   | 80.0 | male   | scalp        |
| 2     | HAM_0002730 | ISIC_0026769 | bkl   | histo   | 80.0 | male   | scalp        |
| 3     | HAM_0002730 | ISIC_0025661 | bkl   | histo   | 80.0 | male   | scalp        |
| 4     | HAM_0001466 | ISIC_0031633 | bkl   | histo   | 75.0 | male   | ear          |
| ...   | ...         | ...          | ...   | ...     | ...  | ...    | ...          |
| 10010 | HAM_0002867 | ISIC_0033084 | akiec | histo   | 40.0 | male   | abdomen      |
| 10011 | HAM_0002867 | ISIC_0033550 | akiec | histo   | 40.0 | male   | abdomen      |
| 10012 | HAM_0002867 | ISIC_0033536 | akiec | histo   | 40.0 | male   | abdomen      |
| 10013 | HAM_0000239 | ISIC_0032854 | akiec | histo   | 80.0 | male   | face         |
| 10014 | HAM_0003521 | ISIC_0032258 | mel   | histo   | 70.0 | female | back         |

10015 rows × 7 columns

In [13]:

```
file.isna().sum()
```

Out[13]:

```
lesion_id      0
image_id       0
dx             0
dx_type        0
age           57
sex            0
localization   0
dtype: int64
```

In [18]:

```
mean_age=int(file['age'].mean())
```

In [19]:

```
file["age"].fillna(mean_age, inplace = True)
```

In [20]:

```
file.isna().sum()
```

Out[20]:

```
lesion_id      0
image_id       0
dx             0
dx_type        0
age            0
sex            0
localization   0
dtype: int64
```

In [24]:

```
print("Number of male cancer patients:",file['sex'].value_counts()['male'])
print("Number of female cancer patients:",file['sex'].value_counts()['female'])
```

```
Number of male cancer patients: 5406
Number of female cancer patients: 4552
```

In [25]:

```
file['dx']
```

Out[25]:

```
0      bkl
1      bkl
2      bkl
3      bkl
4      bkl
...
10010  akiec
10011  akiec
10012  akiec
10013  akiec
10014   mel
Name: dx, Length: 10015, dtype: object
```

In [ ]:

```
file['dx']
```

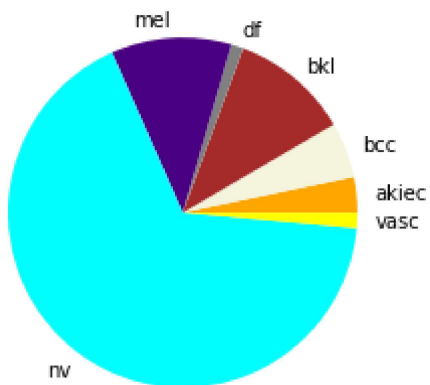
In [34]:

```
dx = file['dx'].value_counts().sort_index()
print(dx)
```

```
akiec    327
bcc      514
bkl     1099
df       115
mel     1113
nv      6705
vasc     142
Name: dx, dtype: int64
```

In [38]:

```
dxtypes=['akiec','bcc','bkl','df','mel','nv','vasc']
colors = ( "orange", "beige", "brown","grey", "indigo", "cyan","yellow")
plt.pie(dx, labels = dxtypes,colors=colors)
plt.show()
```



In [ ]: