### 201700949 설재혁

### 조건부확률 예제 : 한 가족 안의 두 아이의 성별 맞추기

```
In [3]:
```

```
from collections import Counter
import math, random
from IPython.core.interactiveshell import InteractiveShell
InteractiveShell.ast node interactivity = "all"
def random kid():
   return random.choice(["boy", "girl"])
kid test list = [random kid() for i in range(10)]
kid test list #random kid 함수는 boy와 girl 두개의 값중에 하는 램덤하게 추출함
both girls = 0
older girl = 0
either girl = 0
random.seed(0)
for _ in range(10000):
   younger = random kid()
   older = random_kid()
   if older == "girl": # 큰 아이가 여자일 경우 +1
       older_girl += 1
    if older == "girl" and younger == "girl": #둘다 여자일 경우 +1
       both girls += 1
    if older == "girl" or younger == "girl": #둘중에 하나라도 여자일경우 +1
       either girl += 1
print ("P(both | older):", both_girls / older_girl) # 0.514 ~ 1/2 #큰 아이가 딸이고
print ("P(both | either): ", both girls / either girl) # 0.342 ~ 1/3 # 둘중에 한명이
Out[3]:
['girl', 'boy', 'girl', 'girl', 'girl', 'girl', 'boy', 'girl', 'girl',
'girl']
P(both | older): 0.5007089325501317
P(both | either): 0.3311897106109325
```

### 연속분포

#### In [4]:

```
def uniform pdf(x):
    return 1 if x \ge 0 and x < 1 else 0
def uniform cdf(x):
    "returns the probability that a uniform random variable is less than x"
    if x < 0:
                    # uniform random is never less than 0
        return 0
    elif x < 1:
                    # e.g. P(X < 0.4) = 0.4
        return x
    else:
                    # uniform random is always less than 1
        return 1
import numpy as np
x = np.arange(-1.0, 2.0, 0.1)
result array = np.vectorize(uniform cdf, otypes=[np.float])(x)
import matplotlib.pyplot as plt
%pylab inline
plt.plot(x, result array)
plt.axis([-1, 2, -1, 1.5])
plt.show()
```

Populating the interactive namespace from numpy and matplotlib

/Users/seoljaehyeok/opt/anaconda3/lib/python3.8/site-packages/IPython/core/magics/pylab.py:159: UserWarning: pylab import has clobbered thes e variables: ['random'] 
`%matplotlib` prevents importing \* from pylab and numpy

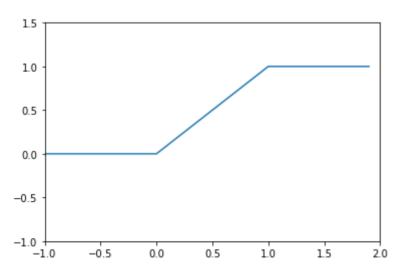
\*matplotlib prevents importing \* from pylab and numpy
warn("pylab import has clobbered these variables: %s" % clobbered +

#### Out[4]:

[<matplotlib.lines.Line2D at 0x7fb8db447df0>]

#### Out[4]:

```
(-1.0, 2.0, -1.0, 1.5)
```



## 정규분포

```
In [5]:
```

```
def normal pdf(x, mu=0, sigma=1):
    sqrt_two_pi = math.sqrt(2 * math.pi)
    return (math.exp(-(x-mu) ** 2 / 2 / sigma ** 2) / (sqrt two pi * sigma))
for sigma value in [1,2,0.5,1]:
    x = np.arange(-6.0, 6.0, 0.1)
    result array = np.vectorize(normal pdf, otypes=[np.float])(x, sigma=sigma value)
      plt.plot(x, result array, "ro")
    plt.plot(x, result array)
plt.axis([-6, 6, 0, 1])
plt.show()
def plot normal pdfs(plt):
    xs = [x / 10.0 \text{ for } x \text{ in } range(-50, 50)]
    plt.plot(xs,[normal_pdf(x,sigma=1) for x in xs],'-',label='mu=0,sigma=1')
    plt.plot(xs,[normal_pdf(x,sigma=0.5) for x in xs],':',label='mu=0,sigma=0.5')
    plt.plot(xs,[normal pdf(x,mu=-1) for x in xs],'-.',label='mu=-1,sigma=1')
    plt.legend()
    plt.show()
import matplotlib.pyplot as plt
plot normal pdfs(plt)
Out[5]:
```

[<matplotlib.lines.Line2D at 0x7fb8db561d60>]

#### Out[5]:

[<matplotlib.lines.Line2D at 0x7fb8db579370>]

#### Out[5]:

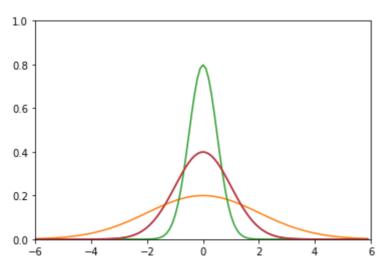
[<matplotlib.lines.Line2D at 0x7fb8db579910>]

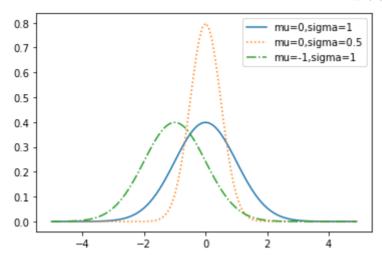
#### Out[5]:

[<matplotlib.lines.Line2D at 0x7fb8db579d90>]

#### Out[5]:

```
(-6.0, 6.0, 0.0, 1.0)
```





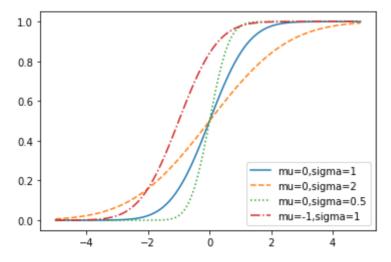
### 표준정규분포

### In [7]:

```
def normal_cdf(x, mu=0,sigma=1):
    return (1 + math.erf((x - mu) / math.sqrt(2) / sigma)) / 2

def plot_normal_cdfs(plt):
    xs = [x / 10.0 for x in range(-50, 50)]
    plt.plot(xs,[normal_cdf(x,sigma=1) for x in xs],'-',label='mu=0,sigma=1')
    plt.plot(xs,[normal_cdf(x,sigma=2) for x in xs],'--',label='mu=0,sigma=2')
    plt.plot(xs,[normal_cdf(x,sigma=0.5) for x in xs],':',label='mu=0,sigma=0.5')
    plt.plot(xs,[normal_cdf(x,mu=-1) for x in xs],'--',label='mu=-1,sigma=1')
    plt.legend(loc=4) # bottom right
    plt.show()

import matplotlib.pyplot as plt
plot_normal_cdfs(plt)
```



### 정규분포 누적 분포 함수의 역함수

```
In [8]:
```

```
def normal cdf(x, mu=0, sigma=1):
    return (1 + math.erf((x - mu) / math.sqrt(2) / sigma)) / 2
def inverse normal cdf(p, mu=0, sigma=1, tolerance=0.00001):
    """find approximate inverse using binary search"""
    # if not standard, compute standard and rescale
    if mu != 0 or sigma != 1:
        return mu + sigma * inverse normal cdf(p, tolerance=tolerance)
    low z, low p = -10.0, 0
                                       # normal cdf(-10) is (very close to) 0
    hi z, hi p = 10.0, 1
                                           # normal cdf(10) is (very close to) 1
    while hi_z - low_z > tolerance:
                                     # consider the midpoint
        mid z = (low z + hi z) / 2
        mid p = normal cdf(mid z) # and the cdf's value there
        if mid p < p:</pre>
            # midpoint is still too low, search above it
            low z, low p = mid z, mid p
        elif mid p > p:
            # midpoint is still too high, search below it
           hi z, hi p = mid z, mid p
        else:
            break
    return mid z
np.vectorize(inverse normal cdf, otypes=[np.float])([0, 0.5, 0.90, 0.95, 0.975, 1])
# 0%, 50%, 90%, 95%, 97.5%, 100%의 확률일경우 누적분포의 확률변수값
Out[8]:
array([-8.75
                  , 0.
                               , 1.28155708, 1.64484978, 1.9599628
```

```
array([-8.75 , 0. , 1.28155708, 1.64484978, 1.9599628
4,
8.75 ])
```

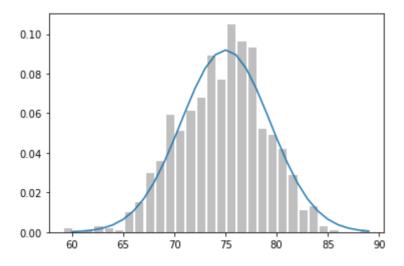
### 중심 극한 정리

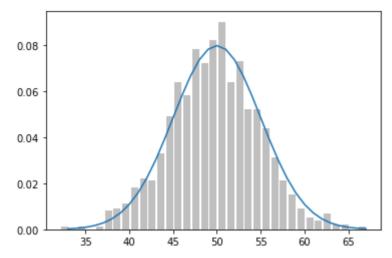
#### In [10]:

```
def bernoulli trial(p):
    return 1 if random.random() 
def binomial(p, n):
    return sum(bernoulli trial(p) for in range(n))
def make hist(p, n, num points):
    data = [binomial(p, n) for in range(num points)]
    # use a bar chart to show the actual binomial samples
    histogram = Counter(data)
    plt.bar([x - 0.4 for x in histogram.keys()],
            [v / num points for v in histogram.values()],
            0.8.
            color='0.75')
    mu = p * n
    sigma = math.sqrt(n * p * (1 - p))
    # use a line chart to show the normal approximation
    xs = range(min(data), max(data) + 1)
    ys = [normal \ cdf(i + 0.5, \ mu, \ sigma) - normal \ cdf(i - 0.5, \ mu, \ sigma)]
          for i in xs]
    plt.plot(xs,ys)
    plt.show()
make hist(0.75, 100, 1000)
make hist(0.50, 100, 1000)
```

<ipython-input-10-dala47944c66>:5: DeprecationWarning: Calling np.sum
(generator) is deprecated, and in the future will give a different res
ult. Use np.sum(np.fromiter(generator)) or the python sum builtin inst
ead.

return sum(bernoulli\_trial(p) for \_ in range(n))





# 201700949 설재혁

In [ ]: