

# Textavinnsla

Strengir: notaðir til að vinna með texta

Gildi: runa af Unicode stöfum

## API

`public class String` (Java string data type)

---

<code>String</code>	<code>String(s)</code>	<i>create a string with the same value as s</i>
<code>int</code>	<code>length()</code>	<i>string length</i>
<code>char</code>	<code>charAt(int i)</code>	<i>i<sup>th</sup> character</i>
<code>String</code>	<code>substring(int i, int j)</code>	<i>i<sup>th</sup> through (j-1)<sup>st</sup> characters</i>
<code>boolean</code>	<code>contains(String sub)</code>	<i>does string contain sub as a substring?</i>
<code>boolean</code>	<code>startsWith(String pre)</code>	<i>does string start with pre?</i>
<code>boolean</code>	<code>endsWith(String post)</code>	<i>does string end with post?</i>
<code>int</code>	<code>indexOf(String p)</code>	<i>index of first occurrence of p</i>
<code>int</code>	<code>indexOf(String p, int i)</code>	<i>index of first occurrence of p after i</i>
<code>String</code>	<code>concat(String t)</code>	<i>this string with t appended</i>
<code>int</code>	<code>compareTo(String t)</code>	<i>string comparison</i>
<code>String</code>	<code>replaceAll(String a, String b)</code>	<i>result of changing a's to b's</i>
<code>String[]</code>	<code>split(String delim)</code>	<i>strings between occurrences of delim</i>
<code>boolean</code>	<code>equals(String t)</code>	<i>is this string's value the same as t's?</i>

# Strengjavinnsla

<i>is the string a palindrome?</i>	<pre>public static boolean isPalindrome(String s) {     int N = s.length();     for (int i = 0; i &lt; N/2; i++)         if (s.charAt(i) != s.charAt(N-1-i))             return false;     return true; }</pre>
<i>extract file name and extension from a command-line argument</i>	<pre>String s = args[0]; int dot = s.indexOf("."); String base = s.substring(0, dot); String extension = s.substring(dot + 1, s.length());</pre>
<i>print all lines in standard input that contain a string specified on the command line</i>	<pre>String query = args[0]; while (!StdIn.isEmpty()) {     String s = StdIn.readLine();     if (s.contains(query)) StdOut.println(s); }</pre>
<i>print all the hyperlinks (to educational institutions) in the text file on standard input</i>	<pre>while (!StdIn.isEmpty()) {     String s = StdIn.readString();     if (s.startsWith("http://") &amp;&amp; s.endsWith(".edu"))         StdOut.println(s); }</pre>

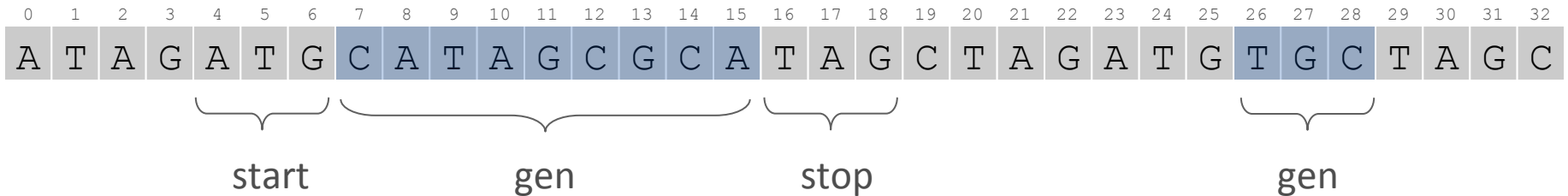
# Genaleit

Erfðamengi: runa af stöfum (A,C,G,T)

Gen: kóði í erfðamenginu sem inniheldur uppskrift af prótíni

- Byrjar á ATG (start codon)
- Margfeldi af 3 kjarnsýrum (3 stafir kóða amínósýru)
- Endar á TAG, TAA eða TGA (stop codon)

Finnum öll möguleg gen í erfðamengi



# Genaleit: reiknirit

## Reiknirit: skönnum vinstri-hægri í erfðamenginu

- Ef við finnum start, setjum beg = i
- Ef við finnum stop og hlutstrengurinn frá beg til i er margfeldi af 3
  - prentum út genið
  - setjum beg = -1

i	codon		beg	gene	<i>remaining portion of input string</i>
	<i>start</i>	<i>stop</i>			
0			-1		ATAGATGCATAGCGCATAGCTAGATGTGCTAGC
1		TAG	-1		ATAGATGCATAGCGCATAGCTAGATGTGCTAGC
4	ATG		4		ATAGATGCATAGCGCATAGCTAGATGTGCTAGC
9		TAG	4		ATAGATGCATAGCGCATAGCTAGATGTGCTAGC
16		TAG	4	CATAGCGCA	ATAGATGCATAGCGCATAGCTAGATGTGCTAGC
20		TAG	-1		ATAGATGCATAGCGCATAGCTAGATGTGCTAGC
23	ATG		23		ATAGATGCATAGCGCATAGCTAGATGTGCTAGC
29		TAG	23	TGC	ATAGATGCATAGCGCATAGCTAGATGTGCTAGC

# Genaleit: útfærsla

```
public class GeneFind {
    public static void main(String[] args) {
        String start = args[0];
        String stop = args[1];
        String genome = StdIn.readAll();

        int beg = -1;
        for (int i = 0; i < genome.length() - 2; i++) {
            String codon = genome.substring(i, i+3);
            if (codon.equals(start)) beg = i;
            if (codon.equals(stop) && beg != -1) {
                String gene = genome.substring(beg+3, i);
                if (gene.length() % 3 == 0) {
                    StdOut.println(gene);
                    beg = -1;
                }
            }
        }
    }
}
```

```
% more genomeTiny.txt
ATAGATGCATAGCGCATAGCTAGATGTGCTAGC

% java GeneFind ATG TAG < genomeTiny.txt
CATAGCGCA
TGC
```

# Strengir og minni

## Strengir eru geymdir í minni sem fylki af stöfum

- `genome = "aacaagtttacaagc"`

D0	D1	D2	D3	D4	D5	D6	D7	D8	D9	DA	DB	DC	DD	DE
a	a	c	a	a	g	t	t	t	a	c	a	a	g	c

genome

A0	A1
D0	15



minnissvæði lengd

- `s = genome.substring(1, 5);`
- `t = genome.substring(9, 13);`

B0	B1	B2	B3
D1	4	D9	4

*s og t vísa á ólíka strengi með sama gildi "acaa"*

- `(s==t)` er false, en `(s.equals(t))` er true

*ber saman tilvísun  
(þ.e. minnissvæði)*

*ber saman stafina í strengjunum*

## Hlutir og klasar - samantekt

- Klasar í Java skilgreina ný gagnatög
- Hlutir í Java eru tilvik af klösum
- Breytur vísa á hluti (ólíkt frumstæðum gagnatögum, eins og með fylki)  
sjá bls. 352-358
- `a == b` er saman tilvísanir `a.equals(b)`  
(ef `a` hefur `equals` aðferð) ber saman  
gildin í hlutunum

# Ný gagnatög

## Til að búa til nýtt gagnatag, skilgreinum við

- Mengi gilda
- Aðgerðir skilgreindar á gildum

## Java klasar: skilgreina gagnatag með

- Tilviksbreytu (instance variable) – mengi gilda
- Aðferð (methods) – aðgerðir á gildum
- Smið (constructor) – býr til og upphafsstillir hlut



# Hleðslu gagnatag

Búum til gagnatag fyrir rafhleðslu í punkti.

Mengi gilda: Þrjár rauntölur ( $x, y$  staðsetning og rafhleðsla)

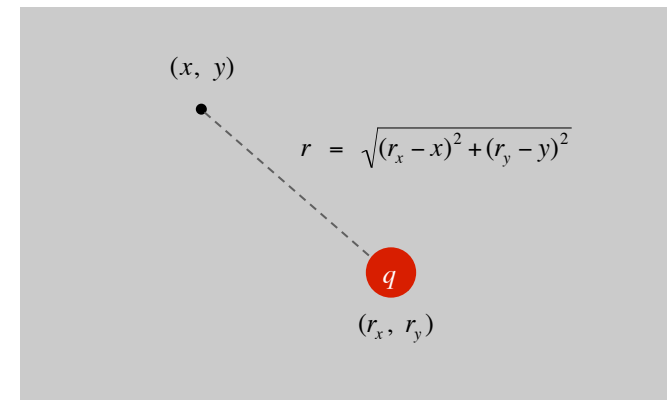
Aðgerðir:

- Búa til nýja hleðslu á  $(r_x, r_y)$  með rafhleðslu  $q$
- Finna spennu í punkt  $(a, b)$  vegna hleðslunnar
- Breyta yfir í streng

$$V = k \frac{q}{r}$$

$r$  = fjarlægð milli  $(x, y)$  og  $(r_x, r_y)$

$k$  = Kúlombfasti =  $8.99 \times 10^9 \text{ N} \cdot \text{m}^2 / \text{C}^2$



Varúð

Ekki hafa áhyggjur af eðlisfræðinni

Það þarf ekki að skilja formúlurnar

Bara breyta þeim í kóða

# Hleðslu gagnatag

Búum til gagnatag fyrir rafhleðslu í punkti.

Mengi gilda: Þrjár rauntölur (x,y staðsetning og rafhleðsla)

## API

```
public class Charge
```

---

```
    Charge(double x0, double y0, double q0)
```

```
    double potentialAt(double x, double y) electric potential at (x, y) due to charge
```

```
    String toString() string representation
```

# Notkun

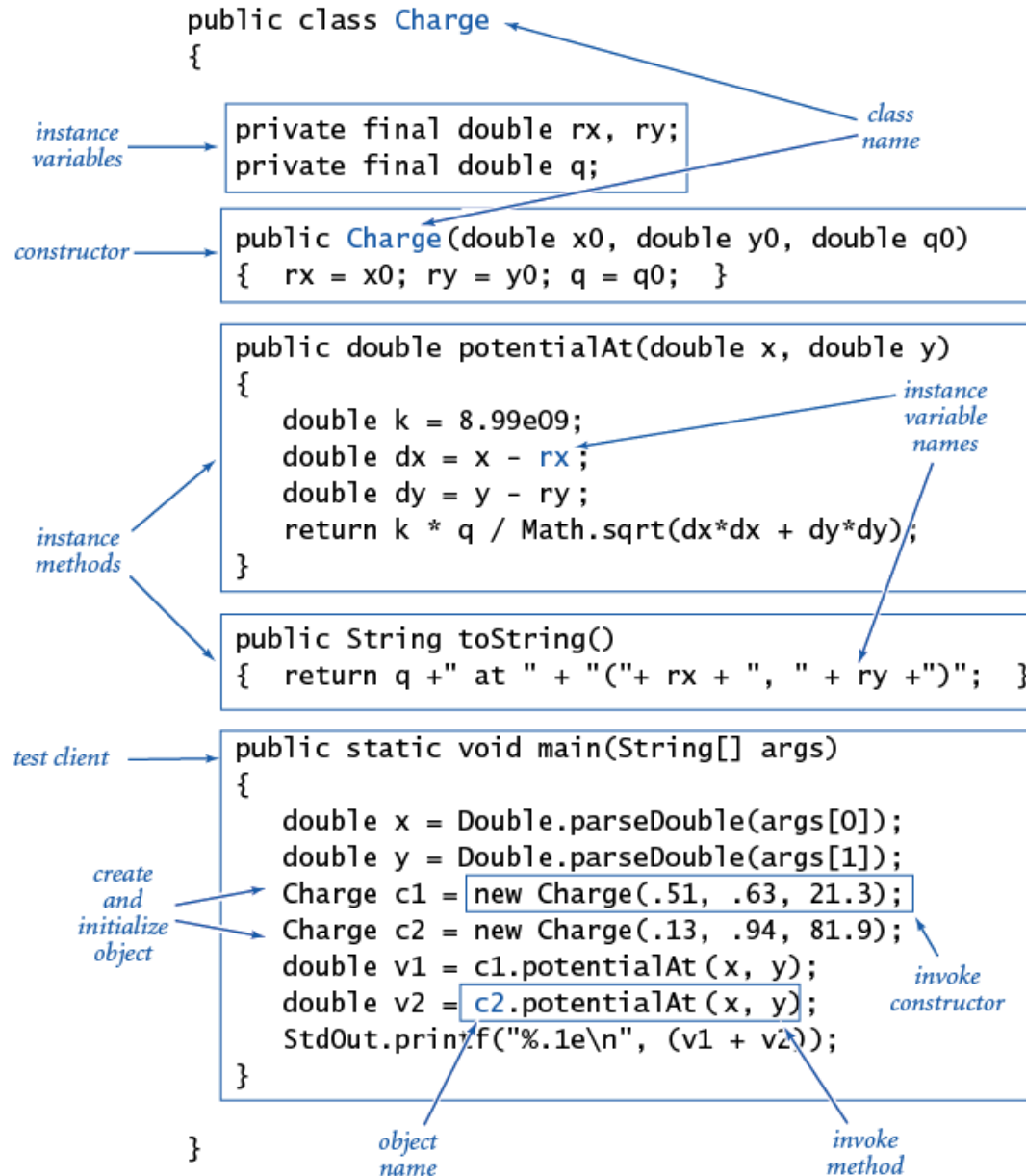
## Prufuforrit: notar klasann og prufar aðferðir

```
public static void main(String[] args) {  
    double x = Double.parseDouble(args[0]);  
    double y = Double.parseDouble(args[1]);  
    Charge c1 = new Charge(.51, .63, 21.3);  
    Charge c2 = new Charge(.13, .94, 81.9);  
    double v1 = c1.potentialAt(x, y);  
    double v2 = c2.potentialAt(x, y);  
    StdOut.println(c1);  
    StdOut.println(c2);  
    StdOut.println(v1 + v2);  
}
```

← Kallar á toString()  
← aðferðina

```
% java Charge .50 .50  
21.3 at (0.51, 0.63)  
81.9 at (0.13, 0.94)  
2.74936907085912e12
```

# Yfirlit



# Tilviksbreytur

## Tilviksbreytur: skilgreina mengi gilda

- Skilgreinum í klasa fyrir utan aðferðir
- Notum alltaf `private` aðgangsheimild
- Notum `final` breytinn fyrir tilviksbreytur sem breytast aldrei

```
public class Charge
{
    private final double rx, ry;
    private final double q;
    .
    .
    .
}
```

The diagram shows a Java class definition for `Charge`. The code is as follows:

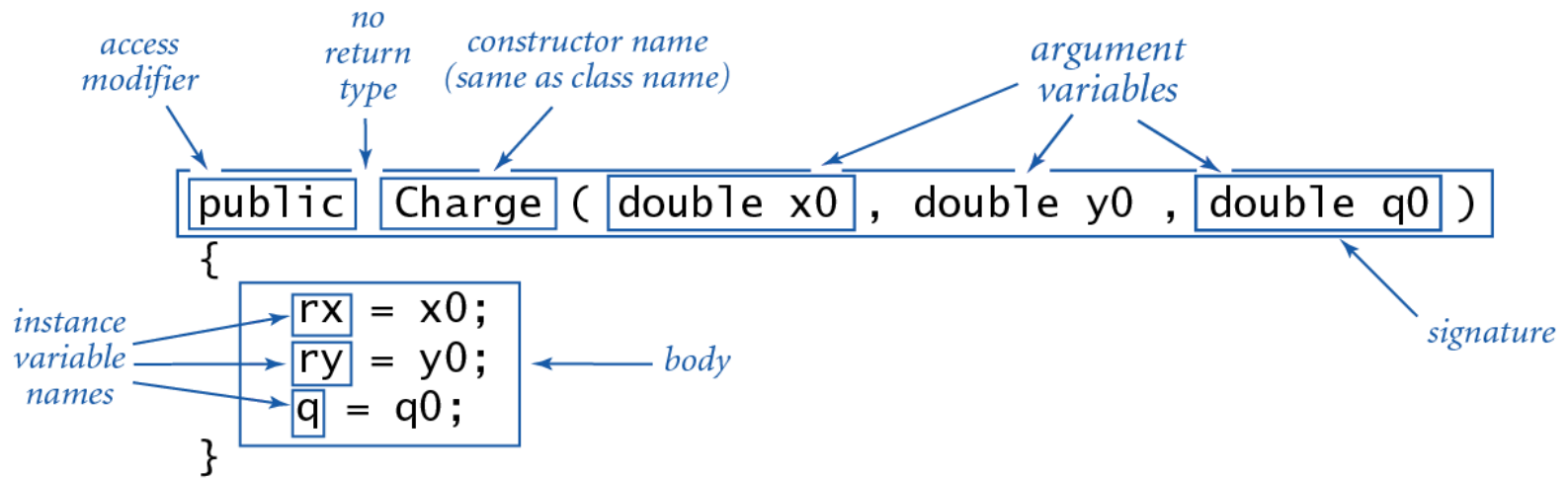
```
public class Charge
{
    private final double rx, ry;
    private final double q;
    .
    .
    .
}
```

Annotations in the diagram:

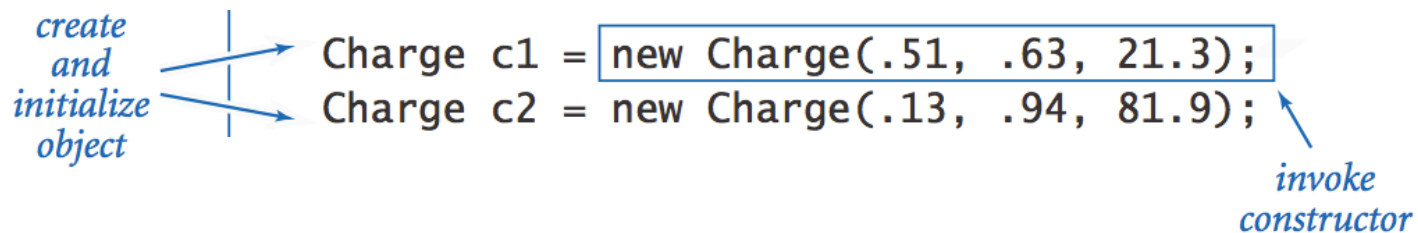
- instance variable declarations*: Points to the two lines of instance variable declarations: `private final double rx, ry;` and `private final double q;`.
- modifiers*: Points to the `private` and `final` keywords in both instance variable declarations.
- public class Charge*: Points to the class declaration line.

# Smiður

Smiður: skilgreinir hvað gerist þegar við búum til hlut

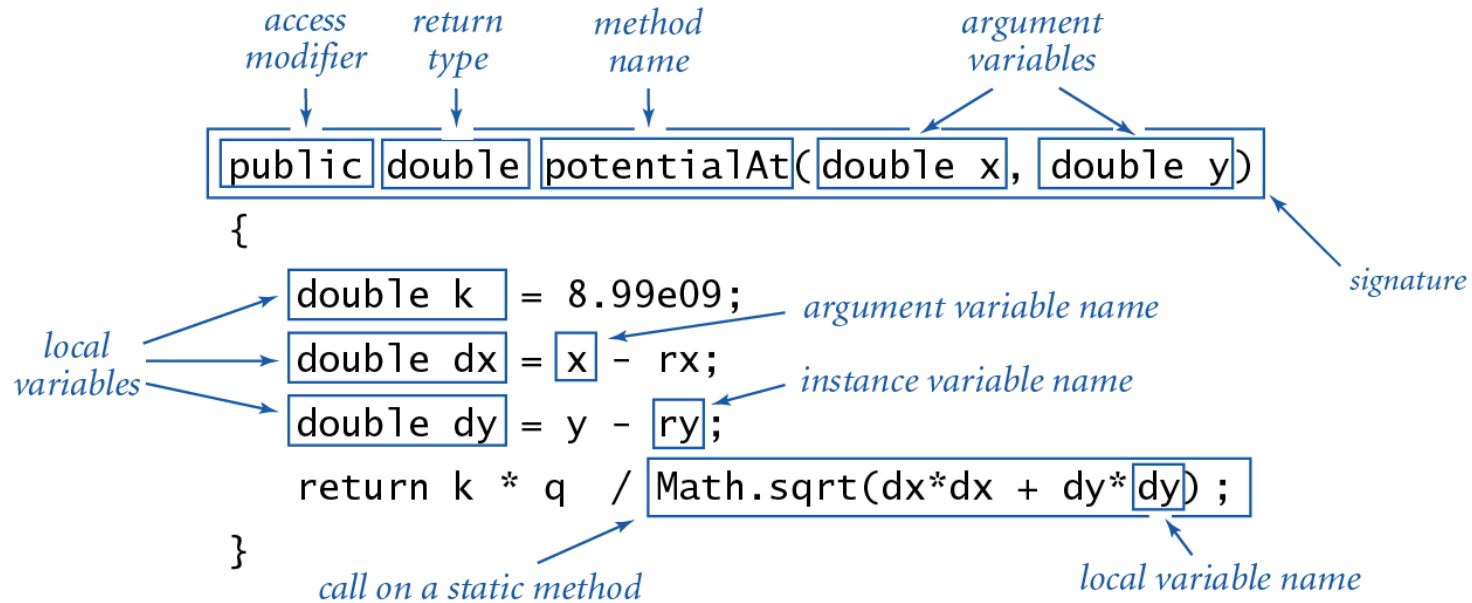


Kallað á smið: notum `new` til að búa til nýjan hlut



# Aðferð

Aðferð: skilgreinir aðgerðir á hlut (tilviksbreytum)



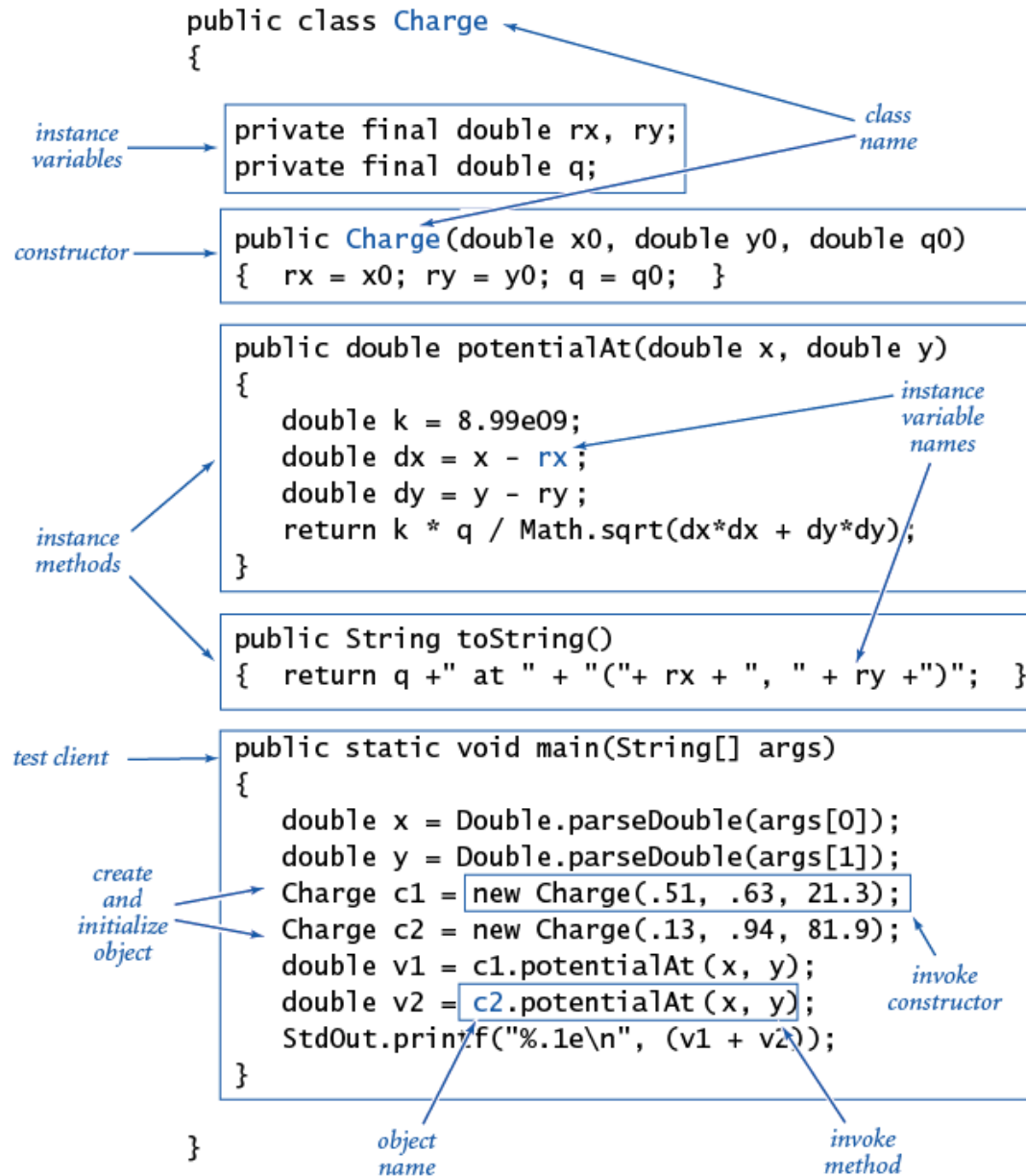
Notkun á aðferð:

```
double v1 = c1.potentialAt(x, y);  
double v2 = c2.potentialAt(x, y);
```

Annotations: *object name* (c2), *invoke method* (c2.potentialAt)



# Geymt í Charge.java



# Birting spennu

Birting spennu: lesum N hleðslur  $(x,y,q)$  af staðal inntaki og reiknum samanlagða spennu í einingaferninginum  $(0,1) \times (0,1)$

```
% more charges.txt
```

```
9
```

```
.51 .63 -100
```

```
.50 .50 40
```

```
.50 .72 10
```

```
.33 .33 5
```

```
.20 .20 -10
```

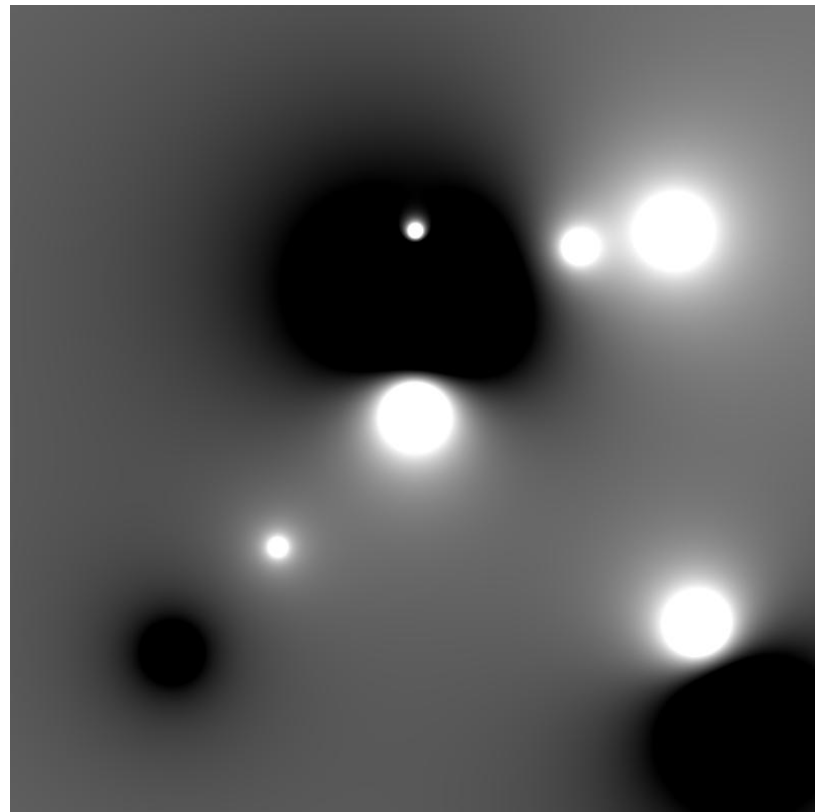
```
.70 .70 10
```

```
.82 .72 20
```

```
.85 .23 30
```

```
.90 .12 -50
```

```
% java Potential < charges.txt
```



# Birting spennu

Höldum utan um allar hleðslurnar í fylki af hlutum.

```
% more charges.txt
9
.51 .63 -100
.50 .50 40
.50 .72 10
.33 .33 5
.20 .20 -10
.70 .70 10
.82 .72 20
.85 .23 30
.90 .12 -50
```

```
// read in the data
int N = StdIn.readInt();
Charge[] a = new Charge[N];
for (int i = 0; i < N; i++) {
    double x0 = StdIn.readDouble();
    double y0 = StdIn.readDouble();
    double q0 = StdIn.readDouble();
    a[i] = new Charge(x0, y0, q0);
}
```

# Birting spennu

```
// plot the data
int SIZE = 512;
Picture pic = new Picture(SIZE, SIZE);
for (int i = 0; i < SIZE; i++) {
    for (int j = 0; j < SIZE; j++) {
        double V = 0.0;
        for (int k = 0; k < N; k++) {
            double x = 1.0 * i / SIZE;
            double y = 1.0 * j / SIZE;
            V += a[k].potentialAt(x, y);
        }
        Color color = getColor(V);
        pic.set(i, SIZE-1-j, color);
    }
}
pic.show();
```

reiknum út lit sem fall af spennu