IMBH Fingerprints in Globular Clusters



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Intermediate Mass Black Holes

- Black Holes of 10²-10⁴ Msun, missing link between stellar and supermassive BHs Have been predicted in different astrophysical scenarios:
 - Remnants of Population III stars (Heger et al. 2003)
 - Runaway Collapse of Young Star Clusters (Portegies-Zwart et al. 2004)
 - Globular Clusters seem the best place to look for them
 - But unambiguous detection is still missing

Searching for IMBHs in GCs

- Globular Clusters have very little gas
 - x-ray emission faint at best
- Sphere of influence of the BH is small (a few arcsecs): Limited Direct BH Influence



- Interpretation of surface brightness + velocity dispersion profiles is model dependent [e.g. through isotropic Jeans Equations]
 - Alternative Dynamic Models with NO BH can be constructed (e.g. Baumgardt et al. 2005)

Searching for IMBHs in GCs

- Proper motion studies can provide the best evidence for IMBH but these are expensive
 - multiyear HST observations needed for GCs
- Are we focusing on the right GCs candidates?
 - Can we identify fingerprints for the IMBH presence?



IMBH fingerprint: rc/rh

Efficient IMBH heating leads to

- Universal large rc/rh after a few relaxation times
- But... there are other (equally) efficient heating sources
- Stellar evolution (Hurley 2007),
 WD kicks (Richer's talk), stellar
 collisions (Chatterjee et al.
 2009), stellar BHs (Davies' Talk)



Trenti et al. (2007)

IMBH fingerprint: mass segregation

- In a GC the most massive stars segregate toward the center of the system (energy equipartition)
- Simulations with an IMBH have less mass segregation (Baumgardt et al. 2004, Trenti et al. 2007)
 - Effect well beyond the BH sphere of influence!



Spatial distribution of binaries @ t=10t_{rh}

Trenti et al. (2007)



Quenching of mass segregation

- IMBH quickly gains at least one tightly bound massive star:
 - A super-scatter machine is born!
- Three body encounters with the BH scatter out incoming stars independently of their mass
- no strong dependence on BH mass expected or seen in simulations when mBH>>mstar
 - random walk of the IMBH within the core: loss cone is constantly replenished, high rate of interactions over time

Our Modeling

- Direct N-body simulations with Aarseth's NBODY6:
 - NO softening
- Exact treatment of all strong interactions including with the BH
 - Up to N=32768
- Grid of initial conditions
 - "Late Time" Mass function, Primordial Binary Fraction, Tidal Field, Concentration
- IMBH mass about 1% of total mass of the system
- Runs carried out until tidal dissolution (about 15 t_{rh})

Measuring Mass Segregation

$\Delta < m > = <m(r = 0) > - <m(r = rh) >$

Mass Segregation $\Delta < m >$ is measured as the difference in average main sequence mass between the center and the half mass radius

Differential measure:

erases dependence on the IMF

Mass not light based:

less sensitive to fluctuations due to small number of giant stars



Mass Segregation Results: Simulations

- Simulations start with no mass segregation
- After about 5 relaxation times equilibrium value of mass segregation is reached
- Good separation of runs with and without an IMBH



Mass Segregation: A first application

- Search for IMBH fingerprint can be applied to well relaxed clusters (t_{rh}<1Gyr)
- Detailed Star Counts are needed, with coverage to at least half-mass radius
- Data and Simulations need to be treated self-consistently
 - e.g. completeness, FOV, measure of structural parameters

NGC 2298



NGC2298 dataset

- **Cluster properties**
 - t_{rh} = 10^{8.41} yr
 - rh = 49″
 - $M_{tot} = 3x10^4 Msun$
- Data Reduction: DeMarchi & Pulone (2007) HST-ACS WFC F606W & F814W --- 10 σ limit @ m₆₀₆=26.5, m₈₁₄=25.0
 - >50%completeness @ 0.2 Msun

NGC 2298



NGC2298: predictions from simulations

- Simulations analyzed between 6 and 8 t_{rh}
- Full radial mass segregation profile has been obtained
- Plot shows 1 and 2σ scatter of the simulated clusters
 - sample of runs (270 snapshots), sample of random projections
 - Good separation IMBH vs NO BH in the center



NGC2298: comparison with simulations

- Observed mass segregation profile is matched very well by simulations
- Cluster is too segregated to be likely to host an IMBH
- Formal limit from the inner two points: >300Msun BH excluded at 3σ CL
- but limiting factor is number of simulations (only 135 snapshots with IMBH)



NGC2298: Error budget

Poisson errors have been estimated by bootstrap (100 synthetic catalogs)

Possible systematic errors from determination of

- Half mass radius. Even a +/- 4" mis-determination only shifts by less than 1σ Poisson error
- Center. We use mass, not light based measure, more stable:
 [0.4" uncertainty at 1σ]
 Miscentering only increases BH rejection confidence level



The future

Larger sample of simulations

- NBODY-6 SSE/GPU code on NCSA Lincoln cluster
 - Improved statistics, wider sampling of initial conditions, larger N
- Suitable HST data are available in the archive for about 15 clusters



Summary

- IMBHs leave multiple fingerprints of their presence in RELAXED globular clusters
- large rc/rh [unfortunately not unique]
- QUENCHING OF MASS SEGREGATION
- Direct N-body simulations show a clear separation in the amount of mass segregation depending on IMBH presence
 - Application to NGC 2298 validates the method
 - no evidence for BH found, limit M_{bh} < 300 Msun at 3σ
 - Analysis of large sample of galactic globular clusters coming soon