The running coupling of many-fermion systems from Monte Carlo Renormalization Group studies

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Fermion - gauge models

- Can be QCD-like: confining, chirally broken

or

- Can have an infrared fixed point: conformal

The most exciting cases are near the emergence of the IRFP

- Just below the theory could be "walking"
- Just above there is a strongly coupled conformal FP

Questions for lattice studies:

- Minimal N_f where the conformal regime develops
- Running of the coupling just below the conformal window
- Properties of the IRFP (anomalous dimension of the mass)

The most interesting questions are frequently the hardest ones







The lattice phase diagram

Lattice simulations can connect the perturbative FP and strong coupling

- Found IRFP? Done 🖌
- No IRFP? Show that it is confining before a bulk transition is reached
- Strong lattice artifacts can interfere



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Monte Carlo Renormalization Group method

- MCRG was designed to study phase the diagram and critical exponents
- It is well suited to connect the lattice weak and strong coupling phases, identify an IRFP and measure critical exponents
- MCRG works in bare coupling space

2-lattice matching method

The method identifies pairs of bare couplings (β, β') where $a(\beta) = a(\beta') / 2$ and predicts the (differential) bare step scaling function $s_b(\beta; 2) = \beta - \beta'$

- At a fixed point $s_b(\beta^*)=0$
 - $s_b(\beta)$ identifies a FP just like the renormalized one!
- The value of \boldsymbol{s}_{b} is related to the scaling dimension of the coupling
 - for AF models $s_b = 3 \ln(2)/(4\pi^2) b_0 + O(g^2)$
 - $s_b > 0$ where the RG β -function is $\beta(g) < 0$ (sorry)
 - s_b in the mass predicts the anomalous dimension of the mass $m = m' 2^{1/y}$



RG flow lines & 2-lattice matching



at a FP with 1 relevant direction:

– do simulations at K and K'

do RG blocking and compare the blocked actions

- if $K^{(n)} = K'^{(n-1)} -> a(K) = a(K')/2$
- the step scaling function is

 $s_b(K)=lim_{n_b \to \infty}$ (K-K')

Every RG transformation should predict the same $s_b(K)$, but

- the location of the FP depends on the RG transformation
- tuning the free parameters in the RG transformation can pull the FP and its RT close, reducing systematical errors



2- lattice matching MCRG - in practice:

Identifies matched couplings (β, β') by comparing expectations values after $n_b (n_b - 1)$ RG blocking steps

- Can be optimized by tuning the free parameter(s) of the RG transformation
- Finite volume effects are largely controlled
- Requires relatively small statistics
 - Has a lot of built-in consistency checks
 - compare several blocking levels
 - compare several operators
 - compare different RG transformations



Renormalization Group transformation

A real space block transformation averages out the short distance modes Many possibilities - I tried 2 types:





Simulations & results

- I use nHYP smeared staggered fermions, N_f flavors in fundamental representation
- Start with well understood models:
 - N_f = 0,4,8 : QCD like
 - $N_f = 16$: conformal with IRFP
 - $N_f = 12$: borderline; needs a new approach
 - \rightarrow Consider different block transformations:
 - Cover deeper range of couplings
 - Might distinguish QCD-like, walikng and conformal behavior



SU(3) pure gauge - test case

The bare step scaling function can be calculated in many ways

- Schrodinger fn; Wilson loop ratios,
- physical observables $r_{0},\,T_{c}$
- RG matching: $32^4 \rightarrow 16^4$ and $16^4 \rightarrow 8^4$



 \bullet Excellent agreement between $r_0,\,T_c$ and MCRG

- Both SF and MCRG approach the perturbative value
- Since at β =6 we can test confinement, we know there is no physical IRFP

SU(3) pure gauge - test case

Compare different RG transformations:



N_f=8 flavors

• Compare different RG transformations:



 N_f =8 is QCD-like, RG flow is governed by the UVFP at g=0

Good agreement between the 3 RG blockings



N_f=16 flavor SU(3) model Do we see a difference?

On the critical surface around an IRFP the flows converge to the FP when $n_b \rightarrow \infty$ With finite n_b the flow picks up the slowest flowing operator



The location of the IRFP depends on The RG transfomation

 $s_{b}(\beta)$ can depend on the blocking (scheme dependent)

This is a signal for non-QCD-like behavior



N_f=16 flavor SU(3) model

Do we see a difference? $16^4 \rightarrow 8^4 \text{ MCRG}$



ORIG blocking shows $s_b(\beta)=0$ around $\beta=7.0$

HYP blocking has an IRFP around β =8.0

HYP2 blocking ($\alpha_{_2}\!,\!\alpha_{_3}$ different) is yet an other IRFP



N_f=12 flavors

Use the same techniques as before; $16^4 \rightarrow 8^4$



- Step scaling function connects to perturbative regime, remains positive
- HYP blocking predicts different $s_b(\beta) \sim 0$
 - Could be scaling violation, but β =5.0-7.0 is very weak coupling

•Could develop an IRFP later

- Could be "walking", the flow is governed by the small value of the β function



N_f=12 flavors

Flow is not (fully) controlled by the perturbative UVFP

- Can one find the IRFP (back flow) or find a confining phase?
 - RG matching dies for β <5.2 (strong lattice artifacts)
 - In the range β = (2.5,5.0) there is no sign for a bulk phase transition
 - 16⁴ lattices are not confined at β =4.5, 8³× 16 lattices are not confined at β =2.5
 - β =2.5 is deep in strong coupling



Conclusion

MCRG is an effective alternative method to study the phase structure and scaling properties of lattice QFT's

- o The method is very universal, straightforward to implement for any other system (sextet fermions are under study)
- o MCRG requires only limited statistics
- o MCRG can predict the anomalous dimension of the mass
- N_f =0-8,16 as expected. N_f =12 is difficult:
 - $s_b(\beta)$ hovers above zero, the flow is extremely slow
 - Different RG transformations predict different flow
 - Flow is not controlled by the perturbative UVFP



MCRG to find the mass anomalous dimension

N_f=16 flavor SU(3) model

Matching in the mass at fixed β = 5.8

 $m_2 = m_1 2^{1/\nu}$



use the same gauge
observables (probably not the best choice)

-at α_{opt} both n_b=2(1) and 3(2) predicts the same matching pair

The critical exponent for the mass

At several couplings, mass values



 $m_2 = m_1 2^{1/\nu}$ $\nu = 1.0(1)$

Free field exponent (close to GFP)

